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Scoring table:
                                                   Sequence:
                                                                Perfect score:
                                                                                   Title:
                                                                                                                                                    Run on:
                                                                                                                                                                                 OM protein -
                                                                                                                                                                                 protein search, using sw model
                                                                  US-09-284-107-21
55
                  BLOSUM62
                                                                                                                              June 4, 2001, 12:11:27; Search time 260.2 Seconds (without alignments)
Gapop 10:0 , Gapext 0.5
                                                   1 VLNASVTSPLLE 12
                                                                                                                                                                                                                               Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                7.418 Million cell updates/sec
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries Minimum DB seq length: 0
Maximum DB seq length: 200000000 Searched: Total number of hits satisfying chosen parameters: 1009251 seqs, 160854530 residues 1009251

Database COMB. pep:

score greater than or equal to the score of the result being and is derived by analysis of the total score distribution is the number of results predicted by chance to have a ter than or equal to the score of the result being printed.

/cgn2_6/ptodata/2/paa/US60_

SUMMARIES

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ç	2,	36	.36	36	36	55	55	55	55	55	Score
	۲. در	65.5	65.5	65.5	65.5	100.0	100.0	100.0	100.0	100.0	· Query Match Length DB
	Sas	292	292	292	104	399	374	261	102	96	ength
ţ	2	18	18	18	15	۲	21	16	23	23	: B
	11C-00-534-835A-315	US-09-483-672A-532	US-09-443-686-532	US-09-429-755-315	US-09-107-532-4325	PCT-US00-05882-1128	US-09-702-021-14	US-09-245-764-7	3 US-60-160-203-4414	US-60-160-209-4252	ID
יייי טלעסוונים אוויי טלע	Company 315 App	Sequence 532, App	Sequence 532, App	Sequence 315, App	Sequence 4325, Ap	Sequence 1128, Ap	Sequence 14, Appl	Sequence 7, Appli	Sequence 4414, Ap	Sequence 4252, Ap	Description

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	45	44	43	42	41	40	39	38	37	36	35	34	ώ ω	32	31	30	29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12	1.
	35	35	ა 5	36	36	36	36	36	36	36	36	36	36	36	36	36	36	36	36	36	36	36	36	36	36	36	36	36	36	36	36	36	36	36	36
	ω.	63.6	ω	5	5	5	5	5	5	65.5	65.5	65.5	65.5	65.5	65.5	65.5	65.5	65.5	65.5	65.5	65.5	65.5	5	65.5	65.5	65.5	65.5	65.5	65.5	65.5	65.5	65.5	65.5	65.5	65.5
	188	149	95	329	329	329	329	329	329	329	329	329	329	329	329	329	329	329	329	329	329	329	329	292	292	292	292	292	292	292	292	292	292	292	292
	16	15	18	22	21	21	21	20	20	20	20	20	20	20	19	19	19	18	18	18	17	16	16	22	21	21	21	20	20	20	20	20	20	19	19
•	-248-796-	-09-107-	-09-417-507-	-09-810-936-	-09-780-669-	US-09-759-143-376	-09-709-729-	9-699-295-	US-09-685-166-376			9-657-279-	-651-	9-636-215-	US-09-593-793A-376	9-536-	325A-29	US-09-483-672A-376	US-09-443-686-376	US-09-429-755-299	US-09-352-616A-376	9-289-198-	9-288-946-	US-09-810-936-315	9-780-	9-759-143-	-09-709-729-	-09-699-295	9-685-166-	9-679-426-	9-657-279	-09-651-236	-09-636-21	-09-593-793A-5	US-09-536-857-532
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	27815,	30497	245	299,	376,	376,	376,	299,	376,	376,	376,	376,	376,	376,	376,	376,	299,	376,	376,	299,	376,	299,	376,	315,	₪	e 532,	e 532,	315,	e 532,	e 532,	e 532,	e 532,	O	532,	Sequence 532, App
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ALIGNMENTS

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NUMBER OF SEQ ID NOS: 4646

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 4252

LENGTH: 96

TYPE: PRT
ORGANISM: HUMAN
FEATURE:
NAME/KEY: VARIANT
LOCATION: (1)...(96)
OTHER INFORMATION: Xaa = Any Amino Acid
US-60-160-209-4252
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                                  δÃ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-60-160-209-4252
                                                                      Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 4252, Application (GENERAL INFORMATION: APPLICANT: BONAZZI, VIVIEN
                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/60/160,209
CURRENT FILING DATE: 1999-10-19
                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: ISOLATED HUMAN KINASE PROTEINS, NUCLEIC TITLE OF INVENTION: ACID MOLECULES ENCODING HUMAN KINASE PROTIENS AND USES TITLE OF INVENTION: THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                 FILE REFERENCE:
7 VLNASVTSPLLE 18
                                  1 VLNASVTSPLLE 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Application US/60160209
                                                                                       Score 55; DB 23
Pred. No. 0.007;
                                                                         Mismatches
                                                                                                           DB 23; Length 96;
                                                                         Indels
                                                                         0;
                                                                         Gaps
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RESULT
                                                                              38-09-702-021-14
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                                                   GENERAL INFORMATION:
                                                                  Sequence 14,
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                                                                                                                                                                                                        Query Match
Best Local Similarity
Matches 12; Conser
                                                                                                                                                                                                                                                                                                                                          EARLIER FILING DATE: 1998-02-06
NUMBER OF SEQ ID NOS: 15
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-60-160-203-4414
   FILE REFERENCE
              APPLICANT: C. Frase
                                                                                                                                                                                                                                                                                     TYPE: PRT ORGANISM: Homo sapiens -09-245-764-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 12; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 6374
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 4414
LENGTH: 102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/245,764
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: THREE DIMENSIONAL STRUCTURES AND MODELS OF FC RECEPTORS FILE REFERENCE: 4102-4
                                                                                                                                                                                                                                                                                                                                                                                                                                              ARLIER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Powell, Maree S. APPLICANT: McKenzie, Ian F.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Hogarth, P. Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/60/160,203
CURRENT FILING DATE: 1999-10-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: BONAZZI, VIVIEN
TITLE OF INVENTION: ISOLATED HUMI
TITLE OF INVENTION: NUCLEIC ACID
TITLE OF INVENTION: USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                         ARLIER APPLICATION NUMBER: 60/073,972
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .PPLICANT: Epa, Vidana
                                                                                                                                                                                                                                                                                                                                   ENGTH: 261
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
                                                                                                                                           174 VLNASVTSPLLE 185
                                                                                                                                                             1 VLNASVTSPLLE 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15 VLNASVTSPLLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 1998-09-1
                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 60/099,994
                                                             Application US/09702021
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Maxwell, Kelly F. Garrett, Thomas P.J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Application US/09245764
                                                                                                                                                                                                     100.0%;
ilarity 100.0%;
Conservative (
                                  Fraser
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N: FAIL MOLECULES AND USES THEREOF 7853-217
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ISOLATED HUMAN SECRETED PROTEINS, NUCLEIC ACID MOLECULES ENCODING HUMAN SECRETED PROTEINS AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%;
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                                                                                                                                                                                                                 Score 55; DB 16
Pred. No. 0.022;
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                                                                                                                                                                                                     Mismatches
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                                                                                                                                                                                                                                Length 261;
                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 102;
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                                                                                                                                                                                                  0,
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                                                                                                                                                                                              Gaps .
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US-09-107-532-4325
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                                                                                                                                                                                                                                                                                                                                                                 ; NAME/KEY: SITE; LOCATION: (349); OTHER INFORMATION: PCT-US00-05882-1128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
"atches 12; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; ORGANISM: Homo sapiens US-09-702-021-14
                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn
SEQ ID NO 1128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 1128,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 14
                                                                                                                                               equence 4325, Application US/09107532 ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: PCT/US00/05882
CURRENT FILING DATE: 2000-03-08
EARLIER APPLICATION NUMBER: 60/124,270
EARLIER FILING DATE: 1999-03-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Steve Ruben
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/702,021
CURRENT FILING DATE: 2000-10-30
NUMBER OF SEQ ID NOS: 45
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
NAME/KEY: SITE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 37
                                                                                                                                                                                                                                                                                                                                                                                                                                       LOCATION: (208)
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TILE OF INVENTION: Human Cancer Associated Gene Sequences and Polypeptides TLE REFERENCE: PA106PCT
                                                         NUMBER OF SEQUENCES: 7
                                                                                                                                   APPLICANT:
                                                                                                      TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ENGTH:
                                                                                        TITLE OF INVENTION:
                                                                                                                                                                                                                                    218 VLNASVTSPLLE
                                                                                                                                                                                                                                                                                                           Local
                                STREET:
                                              ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           193 VLNASVTSPLLE 204
                                                                                                                                                                                                                                                               1 VLNASVTSPLLE 12
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                                                                                                                                                                                                                                                                                                          Similarity
Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Craig Rosen,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FastSEQ for Windows Version 3.0
                          : GENOME THERAPEUTICS CORPORATION 100 Beaver Street
                                                                                              Lynn A Doucette-Stamm and David Bush
"ENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO
"ENTION: ENTEROCOCCUS FAECIUM FOR
                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ID NOS: 1694
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Application
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                                                                                                                                                                                                                                    229
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ver.
                                                                                                                                                                                                                                                                                                                                                                         Xaa equals any of the naturally
                                                                                                                                                                                                                                                                                                                                                                                                                               Xaa equals any of the naturally occurring L-amino acids
                                                                        DIAGNOSTICS AND 7308
                                                                                                                                                                                                                                                                                                        100.0%;
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                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                       Score 55; DE
Pred. No. 0.0
); Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 55; Pred. No. 0
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                                                                                                                                                                                                                                                                                                     DB 1;
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                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                         L-amino acids
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                                                                                                                                                                                                                                                                                      0;
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COMPUTER READABLE FORM:
MEDIUM TYPE: CD/ROM ISO9660

TELEFAX: (781)893-8277 NFORMATION FOR SEQ ID NO:

4325:

ELECOMMUNICATION INFORMATION:

(781)893-5007

TELEPHONE:

REGISTRATION NUMBER: 40,489
REFERENCE/DOCKET NUMBER: GTV

GTC-012

RICE APPLICATION DATA:

APPLICATION NUMBER: 60/051571
FILING DATE: July 2, 1997
TTORNEY/AGENT INFORMATION:
NAME: Ariniello, Pamela Deneke

RIOR APPLICATION DATA:

APPLICATION NUMBER:

60/ 085598

APPLICATION NUMBER: FILING DATE:

US/09/107,532

CURRENT APPLICATION DATA:

SOFTWARE: OPERATING SYSTEM

SEQUENCE CHARACTERISTICS

RIGINAL SOURCE OLECULE TYPE: pro

ropology: ENGTH:

linear

protein

amino acid

104 amino acids

ORGANISM:

Enterococcus faecium

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Query Match
Best Local Similarity bu....
6; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; NAME/KEY: misc_feature
; LOCATION: 1...104
US-09-107-532-4325
                                                                  ORGANISM: Homo sapiens
JS-09-429-755-315
                                                                                                                                                                                                                                                                                                                                           JS-09-429-755-315
                                                                                             SOFTWARE: Fa
SEQ ID NO 315
LENGTH: 292
TYPE: PRT
Query Match 65.
Best Local Similarity 70.
Matches 7; Conservative
                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/429,755A
CURRENT FILING DATE: 1999-10-28
NUMBER OF SEQ ID NOS: 315
                                                                                                                                                                                         ILE REFERENCE:
                                                                                                                                                                                                  ITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                 72 NASITEPFIE 81
                                                                                                                                                                                                                                                                                                                                                                                                                          3 NASVTSPLLE 12
                                                                                                                                                                                                                                                                                                                 INFORMATION
                                                                                                                                     FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                Dillon, Davin
                                                                                                                                                                                                                                            Misher, Lynda
Retter, Marc W
                                                                                                                                                                                                                                                                                      Frudakis, Tony N. Smith, John M.
                                                                                                                                                                                                                                                                         Reed, Steven G.
                                                                                                                                                                                                                                                                                                                             Application US/09429755A
                                                                                                                                                                                           ON: TREATMENT AND DIAGNOSIS 210121.419C6
                                                                                                                                                                                                                   COMPOSITIONS AND METHODS FOR THE
            65.5%;
70.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 65.5%;
Score 36; DB Pred. No. 1.1e. 2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
              1.1e+02;
                          DB 18;
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                                                                                                                                                                                                      OF BREAST CANCER
                        Length 292;
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 Indels
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Gaps
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SEQ ID NO 532
LENGTH: 292
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      밁
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                                              NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ fo
SEQ ID NO 532
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL
                                                                                                   CURRENT APPLICATION NUMBER: US/09/483,672A
CURRENT FILING DATE: 2000-01-14
                                                                                                                                    FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/443,686
CURRENT FILING DATE: 1999-11-18
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                                                                                                                                              PPLICANT: Meagher, Madeleine
ITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
ITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PPLICANT
                                  LENGTH: 292
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND ITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER TILE REFERENCE: 210121.42709A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 259 SASIVSPLLE 268
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity 70. 7; Conservative
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Retter, Mark
Solk, John
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FastSEQ for Windows Version 3.0
                                                                                                                                                                                                               Solk, John A.
Day, Craig H.
Skeiky, Yasir A.W.
                                                                                                                                                                                                                                                                 Kalos, Michael D. Fanger, Gary R. Retter, Marc W.
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Mitcham, Jennifer L.
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                                                                                                                                    210121.42711C11
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                                                                  for Windows Version 3.0
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                                                                                                                                                                                                                                                                                                                                                  Susan Louise
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 36;
Pred. No.
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US-09-483-672A-532

Best Local Similarity
Matches 7; Conserv

Conservative

65.5%;

Score 36; Pred. No.

1.1e+02;

Length 292

Query Match Best Local S Matches

Similarity

65.5%; 70.0%;

Score 36; DB 19; Length 292; Pred. No. 1:1e+02;

Conservative

2; Mismatches

Indels

0;

Gaps

0; Gaps

259 SASIVSPLLE 268

3 NASVTSPLLE 12

09-534-825A-315

Query Match

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NUMBER OF SEQ ID NOS:
SOFTWARE: FESTSEQ for
SEQ ID NO 532
LENGTH: 292
TYPE: PRT
; ORGANISM: Homo sapiens
US-09-536-857-532
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Best Local Similarity /u.v
Thes 7; Conservative
                                                                                                            FILE REFERENCE: 210121.4271
CURRENT APPLICATION NUMBER:
CURRENT FILING DATE: 2000-C
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SOFTWARE: FastSEQ fo:
SEQ ID NO 315
LENGTH: 292
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TYPE: PRT
ORGANISM: Homo sapiens
-09-534-825A-315
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                                                                                                                      PPLICANT: Day, Craig H.
ITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND ITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER INTERPRETATION: 171 / 100 / 126 A ST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/534,825A
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259 SASIVSPLLE 268
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                                                                           FastSEQ for Windows Version 3.0
                                                                                                                                                                                              Stolk, John A. 'Day, Craig H.
                                                                                                                                                                                                                                                             Jiang, Yuqui
Reed, Steven G.
Kalos, Michael D.
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Retter, Marc W.
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Harlocker, Susan L.
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                                                                                                            UMBER: US/09/536,857
2000-03-27
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Pred. No.
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SOFTWARE: FastSEQ for Wi
SEQ ID NO 532
LENGTH: 292
TYPE: PRT
ORGANISM: Homo sapiens
US-09-593-793A-532
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; Sequence 532, Ap
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Best Local Similarity
                                                                                                                                                                                                                 ENERAL
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APPLICANT: Helper, William
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
EILE REFERENCE: 210121.42715C15
CURRENT FILLING DATE: 2000-06-13
CURRENT FILLING DATE: 2000-06-13
CURRENCE SEQ ID NOS: 814
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                                                                                                             PPLICANT
                                                                                                                                                                                                                                    equence 532,
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259 SASIVSPLLE 268
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                                                        Fanger, Gary R. Retter, Marc W. Stolk, John A.
Day, Craig H. Vedvick, Thomas S. Carter, Darrick Li, Samuel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FastSEQ for Windows Version 3.0
                                                                                                                       Jiang, Yuqui
Henderson, F
                                                                                                                                                    Harlocker, Susan L.
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Mitcham, Jennifer L.
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                                                                                                          (alos, Michael D.
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Vedvick, Thomas S.
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Mitcham, Jennifer L.
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                                                                                                                     Robert A.
                                                                                                                                                                                                                                                                                                                                                                                     65.5%;
70.0%;
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                                                                                                                                                                                                                                                                                                                                                                                Score 36; DB 19;
Pred. No. 1.1e+02;
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                                                                                                                                                                                                                                                                                                                                                                                            Length 292
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Query Match
Best Local Similarity
Thes 7; Consery:
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CURRENT APPLICATION NUMBER: US/09/651,236
CURRENT FILING DATE: 2000-08-29
NUMBER OF SEQ ID NOS: 865
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 532
LENGTH: 292
TYPE: PRT
ORGANISM: Homo Sapiens
US-09-651-236-532
RESULT 15
US-09-657-279-532
Sequence 532, Application US/09657279
GENERAL INFORMATION:
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; ORGANISM: Homo sapiens
US-09-636-215-532
                                                                                                                                                                          Query Match 65.9
Best Local Similarity 70.0
Matches 7; Conservative
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SOFTWARE: FastSEQ for
SEQ ID NO 532
LENGTH: 292
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APPLICANT: Hepler, William
IITLE OF INVENTION: COMPOSITI
IITLE OF INVENTION: DIAGNOSIS
FILE REFERENCE: 210121.42717C
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259 SASIVSPLLE 268
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259 SASIVSPLLE 268
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Hepler, William
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Carter, Darrick
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Retter, Marc W.
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illon, Davin C.
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ick, Thomas S.
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210121.42717C17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Robert A.
                                                                                                                                                                          Score 36; DB 20;
Pred. No. 1.1e+02;
2; Mismatches 1
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                                                                                                                                                                                                             Length 292;
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER FILE REFERENCE: 210121.427C19
CURRENT APPLICATION NUMBER: US/09/657,279
CURRENT FILING DATE: 2000-09-06
NUMBER OF SEQ ID NOS: 877
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 532
LENGTH: 292
TYPE: PRT
ORGANISM: Homo sapiens
US-09-657-279-532
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                                                                                                                              Query Match
Best Local Similarity
7; Conserv
Search completed: June Job time: 514 sec
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259 SASIVSPLLE 268
                                                                                                          3 NASVTSPLLE 12
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Kalos, Michael D
Fanger, Gary R.
Retter, Marc W.
Stolk, John A.
Day, Craig H.
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Hepler, William
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Dillon, Davin C.
Mitcham, Jennifer L.
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                                                                                                                                         Conservative
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70.0%;
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                                                                                                                                                       Score 36; DB 20;
Pred. No. 1.1e+02;
                                                                                                                                                                      Length 292;
                                                                                                                                        Indels .
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Perfect.score:
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Maximum DB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score
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length: 2000000000
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Gapop 10.0 , Gapext 0.5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pending_Patents_AA_New:*
/cgn2_6/ptodata/1/paa/PCT_NEW_COMB.pep:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         296725 seqs, 15086421 residues
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Copyright (c) 1993 - 2000 Compugen Ltd
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /cgn2_6/ptodata/1/paa/US06_NEW_COMB.pep:*
/cgn2_6/ptodata/1/paa/US07_NEW_COMB.pep:*
/cgn2_6/ptodata/1/paa/US09_NEW_COMB.pep:*
/cgn2_6/ptodata/1/paa/US09_NEW_COMB.pep:*
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                                 US-09-232-773A-12

US-09-232-773A-6

US-09-232-773A-14

US-09-232-773A-14

US-09-426-371-3

US-09-426-371-3

US-09-829-671-2

US-09-829-671-2

US-09-829-671-2

US-09-270-849B-187805

US-09-270-849B-182930

US-09-270-849B-182930

US-09-270-849B-183902

US-09-270-849B-183902

US-09-270-849B-1813902
                                                                                                                                                                                                                                                                                                               US-09-284-107-21
US-09-590-583-315
US-09-822-827-532
US-09-825-301-4
US-09-439-313-532
US-09-590-583-299
US-09-822-827-376
US-09-825-301-2
US-09-825-301-2
US-09-439-313-376
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US-09-739-449-8794
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Sequence 532, App
Sequence 299, App
Sequence 376, App
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2, Appli
187805,
344, App
182930,
183902,
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117, App
2260, Ap
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3, Appli
181441,
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8401, Ap
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376, App
8794, Ap
10537, A
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21, Applica GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/284,107
CURRENT FILING DATE: 1999-10-25
PRIOR APPLICATION NUMBER: PCT/NL97/00557
PRIOR FILING DATE: 1997-10-07
PRIOR APPLICATION NUMBER: EP 96202791.8
PRIOR FILING DATE: 1996-10-08
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APPLICANT: de Kruif, Cornelis Adridan John
TITLE OF INVENTION: METHODS AND MEANS FOR SELECTING PEPTIDES
TITLE OF INVENTION: AND PROTEINS HAVING SPECIFIC AFFINITY FOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILE REFERENCE: 313632000600
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45	44	43	42	41	40	39	38	37	36	35	34	ယ္	32	31	30	29	28
30	30	30	30	30	30	30	30	30	30	30	30	30	30	30	30	30	30
54.5	54.5	54.5	54.5	54.5	54.5	54.5	54.5	54.5	54.5	54.5	54.5	54.5	54.5	54.5	54.5	54.5	54.5
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PCT-US01-04098A-2979	PCT-US01-04098A-1011	PCT-US00-35017A-1259	US-09-816-494-2	US-09-833-245-1202	PCT-US01-11988-1202	.PCT-US01-04098A-3285	PCT-US01-04098A-1317	US-09-270-849B-182179	PCT-US01-04098A-3640	PCT-US01-04098A-1672	US-09-640-211A-2277	US-09-739-449-10533	US-09-270-849B-181024	US-09-833-245-1199	PCT-US01-11988-1199	US-09-270-849B-183119	US-09-270-849B-192928
Sequence 2979, Ap	Sequence 1011, Ap	Sequence 1259, Ap	Sequence 2, Appli	Sequence 1202, Ap	Sequence 1202, Ap	Sequence 3285, Ap	Sequence 1317, Ap	Sequence 182179,	Sequence 3640, Ap	Sequence 1672, Ap	Sequence 2277, Ap	Sequence 10533, A	Sequence 181024,	Sequence 1199, Ap	Sequence 1199, Ap	Sequence 183119,	Sequence 192928,

ALIGNMENTS

21, Application US/09284107

AFFINITY FOR A TARGET

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; OTHER INFORMATION: Oligopeptides representing the extracellular; OTHER INFORMATION: domain of CD64 US-09-284-107-21
                                                                                                                                                                                                     RESULT
                                                                                                                                                                                     US-09-590-583-315
                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                      Sequence
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Best Local Similarity
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                                                                             APPLICANT:
                                                                                            APPLICANT:
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                              APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 12
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
                                                                                                                                                                                                                                                                  1 VLNASVTSPLLE 12
                                                                                                                                                                                                                                                1 VLNASVTSPLLE 12
T: Wang, Aijun
T: Skeiky, Yasir A.W.
INVENTION: COMPOSITIONS AND METHODS FOR THE
INVENTION: THERAPY AND DIAGNOSIS OF BREAST CANCER
                                                                                     Reed, Steven G.
Smith, John M.
Misher, Linda E
                                                        Misher, Linda E.
Dillon, Davin C.
Retter, Marc W.
                                                                                                                                                                        Application US/09590583
                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                            100.0%;
                                                                                                                                                                                                                                                                                                                            Score 55; DB 5;
Pred. No. 0.00025
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FILE REFERENCE: 210121.419C9

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JS-09-825-301-4
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; ORGANISM: Homo sapiens
US-09-822-827-532
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Best Local Similarity
The 7; Conserve
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US-09-590-583-315
                                              CURRENT APPLICATION NUMBER: US/09/825,301
CURRENT FILLING DATE: 2001-04-02
NUMBER OF SEQ ID NOS: 77
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 4
LENGTH: 292
TYPE: PRT
ORGANISM: Homo sapiens
-09-825-301-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 982
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 532
LENGTH: 292
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Best Local Similarity
Matches 7; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/590,583 CURRENT FILING DATE: 2000-06-08 NUMBER OF SEQ ID NOS: 324 SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 315
                                                                                                                                                     TITLE OF INVENTION: METHOD TITLE OF INVENTION: AND METHOD FILE REFERENCE: 210121.513
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 532, Application:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT FILING DATE: 2001-0
                                                                                                                                                                                                                                                  PPLICANT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
                                                                                                                                                                                                                               PPLICANT
                                                                                                                                                                                                                                                                                                APPLICANT: Houghton, Raymond
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                                                                                                                                                                                                                                                                                                                    INFORMATION
                                                                                                                                                                                                                                                                                                                                 4, Application US/09825301
                                                                                                                                                             T: Zehentner, Barbara
T: Persing, David H.
INVENTION: METHODS, COMPOSITIONS AND KITS FOR THE DETECTION INVENTION: AND MONITORING OF BREAST CANCER
                                                                                                                                                                                                                                                             Molesh, David A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Application US/09822827
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Pred. No.
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Pred. No. 38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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CURRENT APPLICATION NUMBER: US/09/439,313; CURRENT FILING DATE: 1999-11-12; NUMBER OF SEQ ID NOS: 575; SOFTWARE: FastSEQ for Windows Version 3.0; SEQ ID NO 532; LENGTH: 292; TYPE: PRT ORGANISM: Homo sapiens US-09-439-313-532
; ORGANISM: Homo sapien US-09-590-583-299
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Best Local Similarity
7; Conserv
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Best Local Similarity
""+"hes 7; Conserve
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                                                               SEQ ID NO 299
                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                             NUMBER OF SEQ ID NOS: 324
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                 APPLICANT: Skeiky, Yasir A.W.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF BREAST CANCER
FILE REFERENCE: 210121.419C9
CURRENT APPLICATION NUMBER: US/09/590,583
CURRENT APPLICATION NUMBER: US/09/590,583
CURRENT APPLICATION OF SEQ ID NOS: 324
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                                                                                                                                                                                                                                                           PPLICANT:
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                                                                                                                                                                                                                                                                                                                                                               equence 299, Application US/09590583
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                                                ENGTH:
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                                                329
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                                                                                                                                                                                                                                   Misher, Linda E.
Dillon, Davin C.
Retter, Marc W.
                                                                                                                                                                                                                                                                                                          Frudakis, Tony N. Reed, Steven G.
                                                                                                                                                                                                                       Wang, Aijun
                                                                                                                                                                                                                                                                                        Smith, John M.
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ilarity 70.0%;
Conservative
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70.0%;
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Pred. No.
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Pred. No.
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Matches

Local

l Similarity 7; Conserv

Conservative

65.5%; 70.0%;

Score 36; Pred. No.

5;

Length 329;

Mismatches

Indels

0;

Gaps

296 SASIVSPLLE 305

3 NASVTSPLLE 12

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RESULT 9
US-09-439-313-376
; Sequence 376, Application US/09439313
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CURRENT FILING DATE: 2001-03-28
NUMBER OF SEQ ID NOS: 982
SOFTWARE: FRATSEQ for Windows Version 3.0
SEQ ID NO 376
                                                                                                                                                                   Best Local Similarity 70.0 Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 376, Application US/09822827 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                              SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 2 LENGTH: 329
                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                              ORGANISM: Homo sapier -09-825-301-2
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                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/825,301
CURRENT FILING DATE: 2001-04-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER SILE REFERENCE: 210121.534C1
                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 77
                                                                                                                                                                                                                                                                                                                                                                                                                              FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Persing, David H.
TITLE OF INVENTION: METHODS, COMPOSITIONS AND KITS FOR THE DETECTION
TITLE OF INVENTION: AND MONITORING OF BREAST CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Xu, Jiangchun
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TYPE: PRT
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                                                                                                                                                                                                                                                                                                TYPE: PRT
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                                                                                             296 SASIVSPLLE 305
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                                                                                                                                 3 NASVTSPLLE 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Houghton, Raymond L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Zehentner, Barbara
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Jiangchun
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      David A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Davin C.
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70.0%;
                                                                                                                                                                                    Score 36; DB 5;
Pred. No. 43;
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Pred. No.
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RESULT 11
US-09-739-449-10537
Sequence 10537, Application US/09739449
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Best Local Similarity
The 7; Conserv
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                                                                                                                                                                                                                                                                                                                                                       ; ORGANISM: Agrobacterium tumefaciens US-09-739-449-8794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-739-449-8794
                                                                                                                                                                                                                                      Query Match
Best Local Similarity 54...
6; Conservative
                                                                                     GENERAL INFORMATION:
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 376
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SEQ ID NO 8794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 8794, Application US/09739449
          APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
TITLE OF INVENTION: Agrobacterium tumefaciens Genome Sequences and Uses Thereof FILE REFERENCE: 38-10(15490)C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Day, Craig
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C9
CURRENT APPLICATION NUMBER: US/09/439,313
CURRENT FILING DATE: 199-11-12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/739,449
CURRENT FILING DATE: 2000-12-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 09/514,000 PRIOR FILING DATE: 2000-02-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Hinkle, Gregory J.

APPLICANT: Slater, Steven C.

TITLE OF INVENTION: Agrobacterium tumefaciens Genome Sequences and Uses Thereof FILE REFERENCE: 38-10(15490)C.
CURRENT APPLICATION NUMBER: US/09/739,449
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TYPE: PRT
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Reed, Steven G.
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Retter, Mark
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Mitcham, Jennifer L.
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Pred. No.
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Pred. No.
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CURRENT FILING DATE:

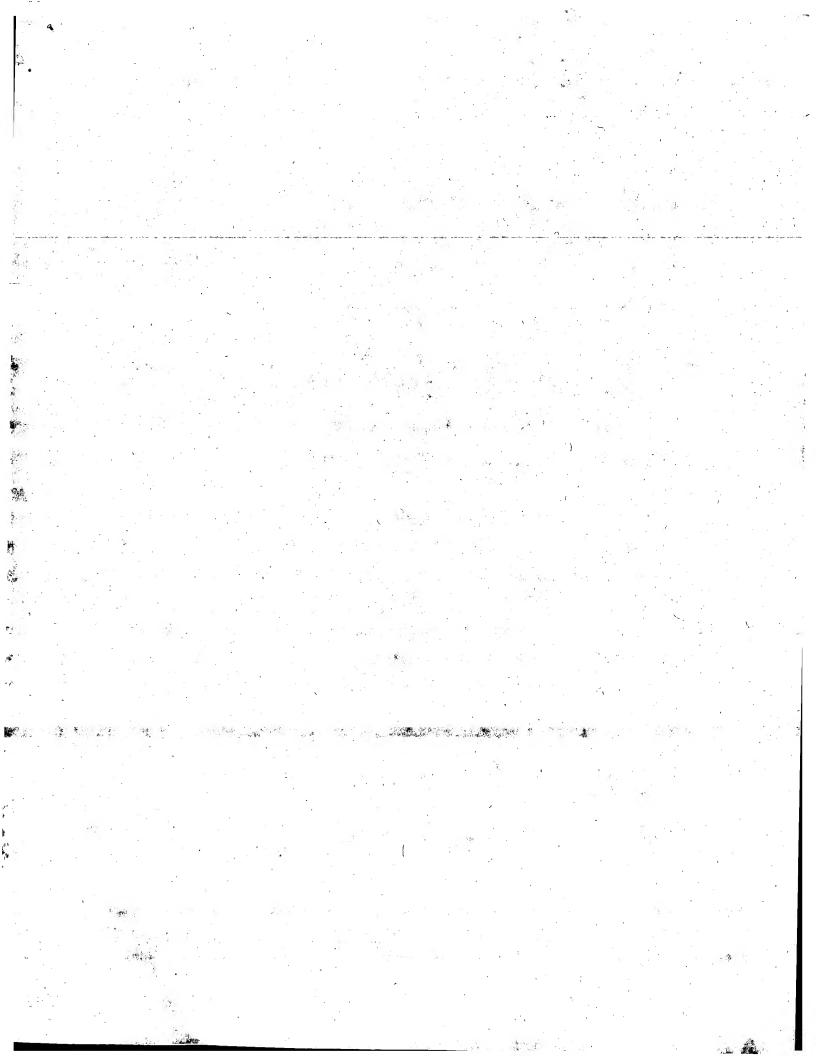
2000-12-19

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; FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Synthetic
US-09-270-849B-194755
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US-09-270-849B-194755
                                                                                                                                                                                                                                                                                                                                                                                             Вb
Query Match
Best Local Similarity bu.v
Thes 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; OTHER INFORMATION: Description of Artificial Sequence: Synthetic; OTHER INFORMATION: at any location having Xaa, Xaa means any amino acid
US-09-270-849B-189198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-270-849B-189198; Sequence 189198, Application US/09270849B; GENERAL INFORMATION:
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CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 195450
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 194755
                                                                                                                                                                                                                                                                                 Sequence 194755, Application US/09270849B GENERAL INFORMATION:
APPLICANT: Swimmer et al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 195450
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 189198
LENGTH.
                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Insect genome survey devices
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Best Local
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PRIOR FILING DATE: 2000-02-23
NUMBER OF SEQ ID NOS: 13351
SEQ ID NO 10537
                                                                                                                                    ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                              FILE REFERENCE:
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CURRENT FILING DATE: 1999-03-17
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                                                                                                                                                                         LENGTH: 341
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ORGANISM: Artificial Sequence
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                                 Score 32;
Pred. No.
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Pred. No. 1.2e+02;
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Pred. No. 73;
                     Mismatches
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                                 2.7e+02;
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                                              Length 341;
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Search completed: June 4, Job time: 590 sec

2001, 12:21:28

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; SOFTWARE: FastSEQ for Wi
; SEQ ID NO 6
; LENGTH: 687
; TYPE: PRT
; ORGANISM: Homo saplen
US-09-232-773A-6
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US-09-232-773A-12
                                        Query Match
Best Local Similarity
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US-09-232-773A-6
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Best Local
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                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/232,773A
CURRENT FILING DATE: 1999-07-15
NUMBER OF SEQ ID NOS: 18
                                                                                                                                                                                                                                                                                                            APPLICANT: Kendall, Richard L. APPLICANT: Thomas, Kenneth A.
                                                                                                                                                                                                                                                                     TITLE OF INVENTION: INHIBITOR OF VASCULAR ENDOTHELIAL CELL TITLE OF INVENTION: GROWTH FACTOR
                                                                                                                                                                                                                                                             FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/232,773A
CURRENT FILING DATE: 1999-07-15
NUMBER OF SEQ ID NOS: 18
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TITLE OF INVENTION: GRA
FILE REFERENCE: 18888DB
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249 VLNCTATTPL 258
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                          1 VLNASVTSPL 10
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                                                      Conservative
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INHIBITOR OF VASCULAR ENDOTHELIAL CELL
GROWTH FACTOR
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                                                                                                                                                                                                  Windows Version 4.0
                                                                  58.2%;
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                                                                 Score 32; I
Pred. No. 5
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Pred. No. 5.4e+02;
                                                        Mismatches
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                                                                    7e+02;
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Result
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
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                                                                                                                                                                                                                                                                                       Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed and is derived by analysis of the total score distribution.
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seq length: 2000000000
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Gapop 10.0 , Gapext 0.5
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61
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Copyright (c) 1993 - 2000 Compugen Ltd.
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                                                                                                                                                                                             Length DB
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                                                                                                                                    US-60-160-209-3193
US-60-160-209-4252
                                                                                                                   US-60-160-203-4414
                                                                                                                                                                                                                                                     SUMMARIES
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7.418 Million cell updates/sec
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Sequence 3133, Ap
Sequence 4252, Ap
Sequence 4414, Ap
Sequence 7, Appli
Sequence 114, Appl
Sequence 1128, Ap
Sequence 10, Appl
Sequence 3220, Ap
Sequence 3255, Ap
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01040	-11501-013/9-7	-US01-01349-5	-US01-01326-	-US01-01326-	-60-197-873-	-60-187-470-	0 - 169 -	-60-147-499-	-09-731-872-	-60-178-307-	-60-173-469-1	-09-528-003-11	us00-29363-13	PCT-US00-07285-116	0	09-380-138-	US01-01349-6	US01-01349-8	-US01-01326-	-60-195-053-182	60-170-374-266	60-194-508-	S-60-170-374-254	07-861-895-2	-09-702-021-3	-09-702-021-3	-09-702-021-	-09-702-021-	-09-702-021-	-09-702-021-	-09-702-021-	-09-702-021-6	-60-185-362-6	-60-185-361-73	115-08-462-973-94
seducates ton other	785	equence 587,	equence 217,	158,	equence 18502	676,	676,		364,		1595,	e 116	134,	116,	e 1,	e 45, App	e 664,		137, Ap	æ	e 2665,	e 3133,	e 2541,	25, A	e 33,	e 32,	31,	30,	eω,	Sequence 5, Appli	34,	e 6, Ap	e 692,	731, A	40

ALIGNMENTS

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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 3193
LENGTH: 58
TYPE: PRT
ORGANISM: HUMAN
US-60-160-209-3193
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                                  RESULT 2
US-60-160-209-4252
Sequence 4252, Application US/60160209 GENERAL INFORMATION:
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Best Local S
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APPLICANT: BONAZZI, VIVIEN
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Sequence 3193, Ap
                                                                                                                                                                                      Matches
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CURRENT FILING DATE: 1999-10-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILE REFERENCE: CL00011
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: ISOLATED HUMAN KINASE PROTEINS, NUCLEIC TITLE OF INVENTION: ACID MOLECULES ENCODING HUMAN KINASE PROTIENS AND USES TITLE OF INVENTION: THEREOF
                                                                                                                                                                                      Local Similarity hes 12; Conserv
                                                                                                                                                1 GNLVTLSCETKL 12
                                                                                                           1 GNLVTLSCETKL 12
                                                                                                                                                                                    100.0%; Score 61; ilarity 100.0%; Pred. No. Conservative 0; Mismatch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Application US/60160209
                                                                                                                                                                                                    No. 0.0021;
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US-60-160-203-4414
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4252
LENGTH: 96
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Matches 12; Conserv
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Best Local :
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SOFTWARE: FastSEQ for
SEQ ID NO 4414
LENGTH: 102
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CURRENT FILING DATE: 1999-10-19
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TITLE OF INVENTION: ISOLATED HUMAN KINASE PROTEINS, NUCLEIC
TITLE OF INVENTION: ACID MOLECULES ENCODING HUMAN KINASE PROTIENS AND USES
TITLE OF INVENTION: THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/60/160,203
CURRENT FILING DATE: 1999-10-19
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LOCATION: (1)...(96)
OTHER INFORMATION: Xaa = Any Amino Acid
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CURRENT APPLICATION NUMBER: US/09/245,764
CURRENT FILING DATE: 1999-02-05
                                                TITLE OF INVENTION: THREE DIMENSIONAL STRUCTURES AND MODELS OF TITLE OF INVENTION: AND USES THEREOF
                                                                                                     APPLICANT: Maxwell, Kelly F. APPLICANT: Garrett, Thomas P.J.
                                                                                                                                         APPLICANT: Hogarth, P. Mark
APPLICANT: Powell, Maree S.
APPLICANT: McKenzie, Ian F.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NERAL INFORMATION:
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                                   FILE REFERENCE: 4102-4
                                                                                        APPLICANT: Epa, Vidana
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                                                                                                                                                                                                                                                                                                       Local Similarity 100
nes 12; Conservative
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Pred. No. 0.0036;
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Pred. No.
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0.0038;
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; ORGANISM: Homo sapiens
US-09-245-764-7
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US-09-702-021-14
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LENGTH: 261
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SEQ ID NO 14
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Best Local Similarity 100
Matches 12; Conservative
                                                                                                                                   NUMBER OF SEQ ID NOS: 1694
SOFTWARE: PatentIn Ver. 2.
SEQ ID NO 1128
LENGTH: 399
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Best Local (
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CURRENT FILING DATE: 2000-10-30
NUMBER OF SEQ ID NOS: 45
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NUMBER OF SEQ ID NOS: 15
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                                                                                                                                                                                                                                                                                                                                                               sequence 1128, Application PC/TUS0005882
                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: PCT/US00/05882
CURRENT FILING DATE: 2000-03-08
EARLIER APPLICATION NUMBER: 60/124,270
EARLIER FILING DATE: 1999-03-12
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                                                                                                                                                                                                                                                                                                              APPLICANT: Craig Rosen, APPLICANT: Steve Ruben
                                                                                                                                                                                                                                                                             FILE REFERENCE: PA106PCT
                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Human Cancer Associated Gene Sequences and Polypeptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 374
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Homo sapiens
                               NAME/KEY: SITE
LOCATION: (208)
OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
                                                                                                    ORGANISM: Homo sapiens
                                                                                    FEATURE:
AME/KEY: SITE
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Llarity 100.0%;
Conservative
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Pred. No.
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Pred. No. 0.014;
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; LOCATION: (349); OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids PCT-US00-05882-1128

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RESULT 9
US-60-170-374-2865
; Sequence 2865, Application US/60170374
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US-09-702-021-10
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SOFTWARE: Fas
SEQ ID NO 10
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Best Local S
Matches 12
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GENERAL INFORMATION:
APPLICANT: C. Fraser
                                                                                                                                 Matches
                                                                                                                                                            Query Match
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TYPE: PRT
ORGANISM: Homo sapiens
-09-702-021-10
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CURRENT FILING DATE: 1999-12-13
NUMBER OF SEQ ID NOS: 4282
                                                                                                                                                                                                                                                                                                                                                                                                                                     60-170-373-3220
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CURRENT FILING DATE: 2000-10-30
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ITLE OF INVENTION:
ITLE OF INVENTION:
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Local Similarity 100.0%;
hes 12; Conservative (
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nes 9; Conserv
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75.08;
                                                                                                                                             78.78;
75.08;
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Pred. No. 0.
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Pred. No.
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SOFTWARE: FastSEQ for Windows Version 4.0; SEQ ID NO 2865; LENGTH: 75; TYPE: PRT; ORGANISM: Human US-60-170-374-2865.
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Best Local Similarity
Matches 9; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/60/170,374
CURRENT FILING DATE: 1999-12-13
NUMBER OF SEQ ID NOS: 3666
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: BODAZZI, VIVIEN
TITLE OF INVENTION: ISOLATE
TITLE OF INVENTION: ACID MOI
TITLE OF INVENTION: THEREOF
FILE REFERENCE: CLOOO146
                                                                                    TELEX: 904136
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
              SEQUENCE CHARACTERISTICS:
LENGTH: 91 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 07/896,457
FILING DATE: 27-MAY 1992
                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Patentin Relication DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
                                                                                                                                     REFERENCE/DOCKET NUMBER: 54
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                                             NAME: BENT, Stephen A. REGISTRATION NUMBER: 2
                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: FILING DATE: 31-007
                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Washington
                                                                                                                      TELEFAX:
                                                                                                                                                                                                                                                                                                      CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE:
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                                                                                                                         (202)672-5399
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HULETT, Mark D.
VENTION: FC-RECEPTOR FOR IMMUNOGLOBULIN EQUENCES: 130
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                                                             PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BONADONNA, Lisa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HOGARTH, Mark P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          McKENZIE, Ian F.C.
linear
                                                                                                                                                                                                                                                                                                                     31-OCT-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bernadette M
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75.0%;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 23;
0.54;
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Length 91;

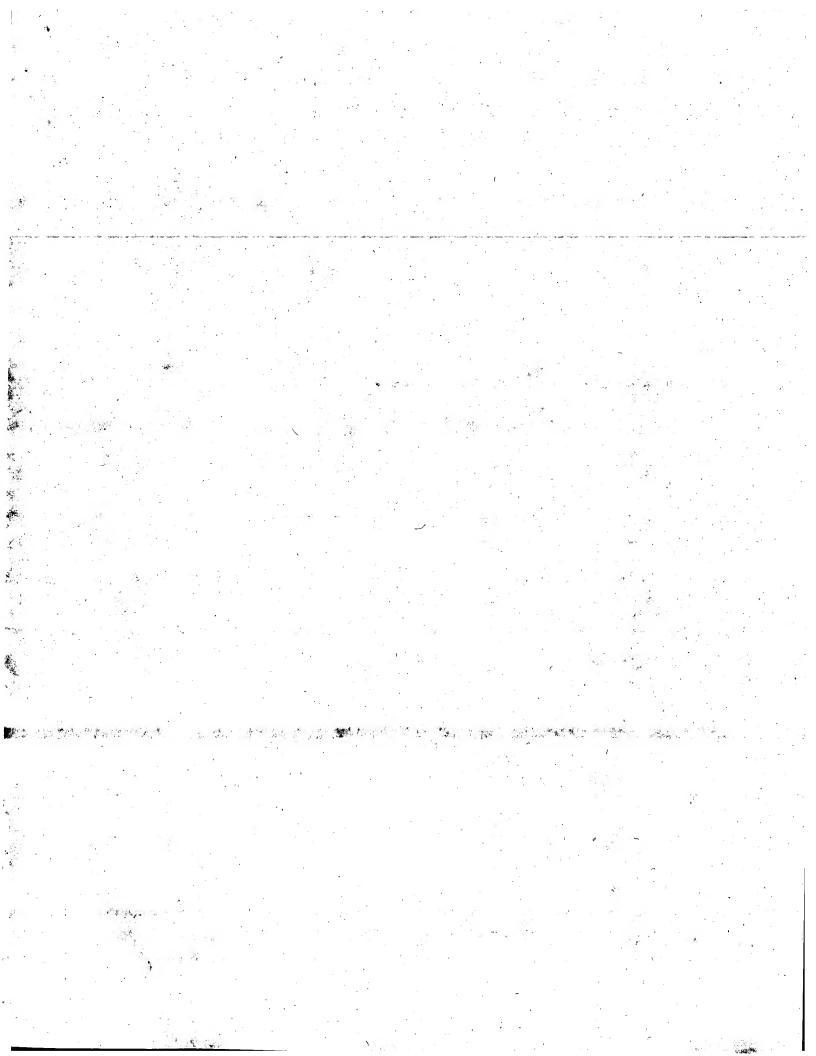
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Query Match
Best Local Similarity
---hes 9; Conserv:
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Best Local
                                                                                                                                                                                                                                                                                                                                                           NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 91 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 1-1904136
                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 16
PIELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
TELEPAX: (202)672-5399
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: VERY PRIOR APPLICATION DATA: APPLICATION NUMBER: VERY PRIOR APPLICATION N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US 0 FILING DATE: 31-OCT-1994 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 26-MAY-1986
TTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
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PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Release #1.0, Version #1.30
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                                                                                                                                                                                                                                       STRANDEDNESS:
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                                                                                                                                                                                                                 TOPOLOGY:
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TITLE OF INVENTION: FC RECEPTOR FOR IMMUNOGLOBULIN
UMBER OF SEQUENCES: 130
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: BENT, Stephen A. REGISTRATION NUMBER: 29,768
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: FILING DATE: 27-MAY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: FILING DATE: 05-JU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HIBBS, Margaret L. SCOTT, Bernadette M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HOGARTH, Mark P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16-SEP-1994
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                                                                                                                                                                                                                                    single
                                                     78.7%;
75.0%;
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75.08;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       us 07/896,
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                  Score 48; DB 8; Lo
Pred. No. 0.66;
2; Mismatches '1;
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Pred. No. 0.66;
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                                                                       Length 91;
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                  Indels
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               0;
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               Gaps
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US-09-702-021-6
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                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
---has 9; Conserv.
                                                                                                                                                                                                                                                                                                                                                                                ; ORGANISM: HUMAN US-60-185-362-692
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US-60-185-362-692
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      В
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                                                                                                  Sequence 6, Application US/09702021 GENERAL INFORMATION:
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APPLICANT: C. Fraser
TITLE OF INVENTION: FAIL MOLECULES AND USES THEREOF
FILE REFERENCE: 7853-217
CURRENT APPLICATION NUMBER: US/09/702,021
CURRENT FILING DATE: 2000-10-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 836
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 692
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 692, Applic GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 731 LENGTH: 177 TYPE: PRT
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/60/185,362
CURRENT FILING DATE: 2000-02-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: ISOLATED HUMAN DRUG TARGET PROTEINS, TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING HUMAN DRUG TARGET PROTEINS, TITLE OF INVENTION: AND USES THEREOF FILE REFERENCE: CL000285
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 731,
                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 177
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Bonazzi, Viven
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/60/185,361
CURRENT FILING DATE: 2000-02-28
NUMBER OF SEQ ID NOS: 968
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: ISOLATED HUMAN SECRETED PROTEINS,
TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING HUMAN SECRETED PROTEINS, AND
TITLE OF INVENTION: USES THEREOF
FILE REFERENCE: CL000291
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2 GSLVTLRCQTKL 13
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75.0%;
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Pred. No. 1
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Page 5
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SEQ ID NO. SEQ ID NOS: 45
SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO. 6
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Title:
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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1. /cgn2_6/ptodata/1/paa/PCT_NEW_COMB.pep:*

2. /cgn2_6/ptodata/1/paa/US06_NEW_COMB.pep:*

3. /cgn2_6/ptodata/1/paa/US07_NEW_COMB.pep:*

4. /cgn2_6/ptodata/1/paa/US08_NEW_COMB.pep:*

5. /cgn2_6/ptodata/1/paa/US09_NEW_COMB.pep:*
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US-09-184-467-20

US-09-184-467-20

US-09-184-467-16

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US-09-184-467-16

US-09-184-467-28

US-60-248-505-847

US-60-248-505-1220

US-09-270-8498-124827

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PCT-US01-11988-1680
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US-09-723-716A-43
US-09-723-716A-68
US-09-723-716A-70
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US-09-270-849B-180519
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133, App
180519,
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US-09-828-825-6	US-09-828-825-16	US-09-828-825-8 .	US-09-828-825-2	US-09-828-825-4	PCT-US00-35017A-1264	PCT-US00-35017A-1153	US-09-832-312-40	US-09-832-312-38	US-09-832-312-36	US-09-832-312-34	US-09-832-312-3	US-09-832-312-5	PCT-US01-10462-13	US-09-832-312-9	US-09-833-245-208	PCT-US01-11988-208 ·	US-08-849-406-38	
Sequence 6, Appli	Sequence 16, Appl	Sequence 8, Appli	Sequence 2, Appli	Sequence 4, Appli	Sequence 1264, Ap	Sequence 1153, Ap	Sequence 40, Appl.	Sequence 38, Appl	Sequence 36, Appl	Sequence 34, Appl	Sequence 3, Appli	Sequence 5, Appli	Sequence 13, Appl	Sequence 9, Appli	Sequence 208, App	Sequence 208; App	Sequence 38, Appl	

ALIGNMENTS

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Query Match
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US-09-284-107-22
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                                                                                                                                                                                                       Sequence 133, Application:
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 22, Applica GENERAL INFORMATION:
                                                        CURRENT APPLICATION NUMBER: PCT/USO1/11988
CURRENT EILING DATE: 2001-01-12
PRIOR APPLICATION NUMBER: 60/229, 358
PRIOR FILING DATE: 2000-04-12
                                                                                                                                          APPLICANT: Human Genome Sciences, Inc.
TITLE OF INVENTION: Albumin Fusion Proteins
FILE REFERENCE: PF546PCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/284,107
CURRENT FILING DATE: 1999-10-25
PRIOR APPLICATION NUMBER: PCT/NL97/00557
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: de Kruif, Cornelis Adriaan John
TITLE OF INVENTION: METHODS AND MEANS FOR SELECTING
TITLE OF INVENTION: AND PROTEINS HAVING SPECIFIC AF
FILE REFERENCE: 313632000600
                   PRIOR APPLICATION NUMBER: 60/256, 931 PRIOR FILING DATE: 2000-12-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Logtenberg, Ton
APPLICANT: de Kruif, Corn
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TYPE: PRT
ORGANISM: Artificial Sequence
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1 GNLVTLSCETKL 12
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60/199, 384
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Pred. No.
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PRIOR FILING DATE: 2000-04-25
NUMBER OF SEQ ID NOS: 2267
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 133

Matches

Conservative

70.0%;

Pred. No.

ed. No. 24; Mismatches

Indels

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Best Local Similarity

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Query Match
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Thes 8; Conserv:
; OTHER INFORMATION: Description of Artificial Sequence: Synthetic US-09-270-849B-180519
                                                                                                                                                                                                                                                                                      RESULT 4
US-09-270-849B-180519
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; ORGANISM: Homo sapiens
US-09-833-245-133
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; ORGANISM: Homo sapiens
PCT-US01-11988-133
                                                                    CURRENT APPLICATION NUMBER: US/09/270,849B
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 195450
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 180519
LENCTH: 257
TYPE: PRT
                                                                                                                                                                                                                                                     Sequence 180519, App
GENERAL INFORMATION:
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TITLE OF INVENTION: Insect genome survey devices
                                                                                                                                                                                                    FILE REFERENCE:
                                  FEATURE:
                                               ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: 60/199, 384
PRIOR FILING DATE: 2000-04-25
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PRIOR FILING DATE: 2000-04-12
PRIOR APPLICATION NUMBER: 60/256, 931
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ITLE OF INVENTION: Albumin Fusion Proteins
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66.7%;
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Query Match

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38;

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; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-565-522B-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-723-716A-43
SEQ ID NO 43
LENGTH: 28
                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 43, Application US/09723716A
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                                                                                            PRIOR APPLICATION NUMBER: GB 9810759.2 PRIOR FILING DATE: 1998-05-19 PRIOR APPLICATION NUMBER: GB 9821129.5
                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/723,716A CURRENT FILING DATE: 2000-11-28 PRIOR APPLICATION NUMBER: US 09/335,087 PRIOR FILING DATE: 1999-06-17 PRIOR APPLICATION NUMBER: PCT/GB99/01588 PRIOR FILING DATE: 1999-05-19
                                     SOFTWARE: PatentIn
                                                          NUMBER OF SEQ ID NOS:
                                                                             PRIOR FILING DATE: 1998-09-29
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                                                                                                                                                                                                                                                                          APPLICANT: WILLCOX, Benjamin Ernest
APPLICANT: BOULTER, Jonathan Michael
TITLE OF INVENTION: Soluble T Cell Receptor
FILE REFERENCE: 102286.499DIV
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                                                                                                                                                                                                                                                                                                                                                                                                                      ENERAL INFORMATION
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PRIOR FILING DATE: 2000-01-21
NUMBER OF SEQ ID NOS: 12
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PRIOR FILING DATE: 2000-04-25
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APPLICANT: Mize, Nancy K
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ITLE OF INVENTION: Methods and Materials Relating to Novel
ITLE OF INVENTION: Polypeptides and Polynucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LE REFERENCE: HYS-8
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                                                                                                                                                                                                                                                                                                                                                       GAO, George Fu
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Liu, Chenghua
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 38;
Pred. No.
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                                                                                                                                                                                                            RESULT
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SOFTWARE: PatentIn vers
SEQ ID NO 68
LENGTH: 287
TYPE: PRT
                                                                                                                                          Sequence 70, Application US/09723716A GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 68, Application US/09723716A
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Best Local Similarity
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PRIOR APPLICATION NUMBER: PCT/GB99/01588
PRIOR FILING DATE: 1999-05-19
PRIOR APPLICATION NUMBER: GB 9810759.2
PRIOR FILING DATE: 1998-05-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               09-723-716A-68
                                                                                                        APPLICANT:
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CURRENT APPLICATION NUMBER: US/09/723,716A
CURRENT FILING DATE: 2000-11-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: JAKOBSEN, Bent Karsten
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                                                                                                                       APPLICANT: JAKOBSEN,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: Description of Artificial Sequence: Amino acid OTHER INFORMATION: sequence of human HLA-A2/flu matrix peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RIOR APPLICATION NUMBER: US 09/335,087
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: Predicted amino acid sequence of the soluble OTHER INFORMATION: HLA-A2/flu matrix restricted beta chain from JM22, OTHER INFORMATION: as fused to the "lecine zipper" domain of c-fos. OTHER INFORMATION: (Figure 11).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Homo sapiens
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T: WILLCOX, Benjamin Ernest
T: BOULTER, Jonathan Michael
INVENTION: Soluble T Cell Receptor
ERENCE: 102286.409DIV
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                                                                              BELL, John Irving
GAO, George Fu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       leucine zipper domain.
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                                                                                                                           Bent Karsten
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66.7%;
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Pred. No.
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Pred. No.
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Sequence 36, Application US/09514009
GENERAL INFORMATION:
APPLICANT: Fraser, Christopher C.
TITLE OF INVENTION: SECRETED PROTEINS AND USES
FILE REFERENCE: 07334-251001
CURRENT APPLICATION UNMERR: US/09/514,009
CURRENT FILING DATE: 2000-02-25
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Best Local Similarity

Matches 7; Conserv
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Best Local Similarity
Matches 8; Conserval
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 46
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CURRENT FILING DATE: 2000-02-25
PRIOR APPLICATION NUMBER: US 09/259,387
PRIOR FILING DATE: 1999-02-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: SECRETED PROTEINS AND USES THEREOF FILE REFERENCE: 07334-251001
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PRIOR FILING DATE: 1998-05-19
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CURRENT FILING DATE: 2000-11-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
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OTHER INFORMATION: Description of Artificial Sequence: Amino acid
OTHER INFORMATION: sequence of human HLA-A2/flu matrix peptide
OTHER INFORMATION: restricted JM22 TCR beta chain fused to c-fos
OTHER INFORMATION: leucine zipper domain and BirA biotinylation tag.
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                                                                                                                                                                                                                                                                                                          57.48;
nilarity 63.68;
Conservative
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                                                                                                                                                                                                                                                                                                                                  Score 35;
Pred. No.
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Pred. No.
                                                                  THEREOF
                                                                                                                                                                                                                                                                                                                                                       Length 57;
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PRIOR APPLICATION NUMBER: US 09/259,387 PRIOR FILING DATE: 1999-02-26

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; NAME/KEY: SIGNAL
; LOCATION: (1)...(34)
US-09-514-009-33
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Best Local Similarity backhes 7; Conservative
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                                                                                                                                                                                                                                                                 S-09-514-009-33
Sequence 33, Application US/09514009
GENERAL INFORMATION:
APPLICANT: Fraser, Christopher C.
TITLE OF INVENTION: SECRETED PROTEINS AND USES THEREOF
FILE REFERENCE: 07334-251001
CURRENT APPLICATION NUMBER: US/09/514,009
CURRENT APPLICATION NUMBER: US/09/514,009
CURRENT APPLICATION NUMBER: 073/760,787
                         Query Match
Best Local Similarity
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Best Local Similarity 63.0
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                                                                                                                                                                       SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 33
LENGTH: 239
TYPE: PRT
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                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 51
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 36
                                                                                                                                               ORGANISM: Mus musculus
                                                                                                                                         FEATURE:
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CURRENT FILING DATE: 2000-02-25
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TITLE OF INVENTION: SECRETED PROTEINS AND USES THEREOF
FILE REFERENCE: 07334-251001
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NUMBER OF SEQ ID NOS: 51
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          Conservative
                         57.4%;
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63.6%;
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63.6%;
                     Score 35; DB 5;
Pred. No. 77;
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Pred. No.
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Pred. No. 37;
1; Mismatches
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                                   Length 239;
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RESULT 14
US-09-184-467-24
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                                                                                                                           Sequence 24, Application US/09184467 GENERAL INFORMATION:
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US-09-184-467-20
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (212) 664-0525
TELEX: 422523 COOPUI
INFORMATION FOR SEQ ID NO: 20:
SEQUENCE CHARACTERISTICS:
LENGTH: 479 amino acids
                                                                                                                                                                                                                                             380 GNTITLPCRIK 390
                                                                                                                                                                                                                                                                               1 GNLVTLSCETK 11
NUMBER OF SEQUENCES: 29 CORRESPONDENCE ADDRESS:
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                                                                    TITLE OF INVENTION: HIV-1 VACCINES,
                                                                                                            APPLICANT: Hasel, Karl W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 08/530,146
FILING DATE: <UNKNOWN>
ATTORNEY/AGENT INFORMATION:
NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 41190/JPW/AJM
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 GNLVTLSCETK 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Release #1.24 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Hasel, Karl W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: HIV-1 VACCINES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/09/184,467 FILING DATE: 02-Nov-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: New York
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Cooper & Dunham
STREET: 30 Rockefeller Plaza
                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                57.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (212)
                                     THERETO, AND THERAPEUTIC THEREOF
                                                                                                                                                                                                                                                                                                                1; Mismatches
                                                                                                                                                                                                                                                                                                                                    Score 35;
Pred. No. 1
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                                               ANTIBODY COMPOSITIONS RELATED ERAPEUTIC AND PROPHYLACTIC USES
                                                                                                                                                                                                                                                                                                                                    5e+02;
                                                                                                                                                                                                                                                                                                            4; Indels
                                                                                                                                                                                                                                                                                                                                                  Length 479;
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ADDRESSEE: Cooper & Dunham STREET: 30 Rockefeller Plaza

CITY: New York STATE: New York COUNTRY: USA

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.24

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/184,467

FILING DATE: 02-Nov-1998

CLASSIFICATION:
 CONTROL OF APPLICATION
 CONTROL OF APPLICA

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RESULT 15
US-09-184-467-16
; Sequence 16, Application US/09184467
; GENERAL INFORMATION:
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; MOLECULE TYPE: protein
; SEQUENCE DESCRIPTION: SEQ ID NO: 24:
US-09-184-467-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 08/530,146

FILING DATE: CURKNOWN>
ATTORNEY/AGENT INFORMATION:

NAME: White, John P.

REGISTRATION NUMBER: 28,678

REFERENCE/DOCKET NUMBER: 41190/JPW/AJM

TELECOMMUNICATION INFORMATION:

TELEPHONE: (212) 977-9550

TELEPAX: (212) 664-0525

TELEX: 422523 COOPUI

INFORMATION FOR SEQ ID NO: 24:

SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 57.4%; Score 35; DB 5; Best Local Similarity 54.5%; Pred. No. 1.5e+02;
COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk.

COMPUTER: IBM.PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.24

CURRENT APPLICATION NUMBER: US/09/184,467

FILING DATE: 02-NOV-1998
CLASSIFICATION LOTA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 GNLVTLSCETK 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: HIV-1 VACCINES, ANTIBODY COMPOSITIONS RELATED THERETO, AND THERAPEUTIC AND PROPHYLACTIC USES
                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Hasel, Karl W
                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6; Conservative
                                                                                                                                                                                                                                                                                 COUNTRY: USA
                                                                                                                                                                                                                                                                                                      CITY: New York
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 479 amino acids TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Cooper & Dunham
STREET: 30 Rockefeller Plaza.
                                                                                                                                                                                                                                                                                                                                                                                                                                               THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1; Mismatches
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В
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Search completed: June 4, 2001, 12:21:28
Job time: 590 sec
                                                                                                                                               Query Match 57.
Best Local Similarity 54.
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                  TELEFAX: (212) 664-0525
TELEX: 422523 COOPUI
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
                                                                                 408 GNTITLPCRIK 418
                                                                                                  1 GNLVTLSCETK 11
                                                                                                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 977-9550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 41190/JPW/AJM
                                                                                                                                                                                                                                                                                                   TYPE: amino acid
                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
                                                                                                                                                                 57.48;
                                                                                                                                                                 Score 35;
Pred. No.
                                                                                                                                                   1; Mismatches
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GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.

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Result
No.
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Maximum DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Title:
Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OM protein - protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Run on:
                                                                                                                                                                                                                                     Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         seq length: 0
seq length: 2000000000
395555561
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                                                                                                                                                                 Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-284-107-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          June 4, 2001, 12:20:01; Search time 260.2 Seconds (without alignments)
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                                                                                                                                                                   Length
                                                                                                                                                                                                                                                                                                   /cgn2_6/ptodata/2/paa/US097_COMB.pep
/cgn2_6/ptodata/2/paa/US098_COMB.pep
/cgn2_6/ptodata/2/paa/US60_COMB.pep:
58
96
102
374
399
91
91
410
261
                                                                                                                                                                   BB
                         US-08-332-562-94
US-08-462-973-94
US-07-861-895-25
                                                                                                                     US-60-160-209-3193
US-60-160-209-4252
                                                                                                                                                                                                              SUMMARIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           7.418 Million cell updates/sec
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                    Sequence 94, Appl
Sequence 94, Appl
Sequence 25, Appl
                                                                                                                                                                Description
                                                                      Sequence 14, Appl
Sequence 1128, Ap
                                                                                                   Sequence 3193, Ap
Sequence 4252, Ap
Sequence 4414, Ap
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	45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26	25	24	23	22	21	20	19	18		16	15	14		12	11
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	23	23	16	23	23	23	23	16	16	17	18	16	14	21	17	16	16	14	14	13	12	11	21	17	13	12	11	23	16	23	23	23	23	23	18
	-60-191-6	-60-191-6:	-09-252-99	-60-191-60	-60-191-6	0-173-4	-60-161-9:	-09-252-69	2-69	-09-328-3	-09-489-0	-09-293-50	-09-060-9394	1-11	-09-398-239-	-09-293-505	93-505	-09-060-93	US-09-060-939A-16	US-08-953-823-2	7-798-	-08-720-484-	-09-711-111-	US-09-398-239-4	-08-953-823-	-08-897-798	-08-720-484-	-60-229-515-7	-09-252-991A	-60-191-681-3	-60-191-637	3-464	-60-167-21	-146-055-7	US-09-489-039A-13963
		Sequence 26977, A	28773,	1814	22997,	18875,	e 2494,	e 7682,	e 7682,	664	e 985	15,	e 15,	7, 1	'n	.17	16,		16,		e 7,	2,	'n	4,	4,	6	4	758	e 17591	æ	477,	e 405,	452,	774, Ap	Sequence 13963, A

ALIGNMENTS

US-60-160-209-4252; Sequence 4252, Application US/60160209; GENERAL INFORMATION: 밁 Qy ; ORGANISM: HUMAN US-60-160-209-3193 RESULT 1 US-60-160-209-3193; Sequence 3193, Ap NUMBER OF SEQ ID NOS: 4646
SOFTWARE: FASTSEQ for Windo
SEQ ID NO 3193
LENGTH: 58
TYPE: PRT Query Match
Best Local Similarity
Matches 12; Conser GENERAL INFORMATION: APPLICANT: BONAZZI, VIVIEN
TITLE OF INVENTION: ISOLATED HUMAN KINASE PROTEINS, NUCLEIC
TITLE OF INVENTION: ACID MOLECULES ENCODING HUMAN KINASE PROTIENS AND USES
TITLE OF INVENTION: THEREOF
FILE REFERENCE: CL000113 CURRENT APPLICATION NUMBER: US/60/160,209
CURRENT FILING DATE: 1999-10-19 13 1 LLQRPGLQLYFS 12 LLQRPGLQLYFS Application US/60160209 Conservative 24 Windows Version 4.0 100.0%; Score 61; DB 23 Pred. No. 0.001; Mismatches DB 23; Length 58; 0 0

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US-09-702-021-14

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US-09-702-021-14
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US-60-160-203-4414
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; ORGANISM: HUMAN
JS-60-160-203-4414
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: VARIANT
LOCATION: (1)...(96)
OTHER INFORMATION: Xaa = Any Amino Acid
US-60-160-209-4252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO 4414
LENGTH: 102
TYPE: PRT
                                                        NUMBER OF SEQ ID NOS: 45
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 14
                                                                                                                                                                                                      Sequence 14, Application US/09702021 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                         Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 12; Conservative (
                                                                                                          CURRENT APPLICATION NUMBER: US/09/702,021 CURRENT FILING DATE: 2000-10-30
                                                                                                                                      APPLICANT: C. Fraser
TITLE OF INVENTION: FAIL MOLECULES AND USES THEREOF
FILE REFERENCE: 7853-217
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 4414, Application US/60160203 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 12; Conserv
                         TYPE: PR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 6374
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/60/160,203
CURRENT FILING DATE: 1999-10-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ ID NO 4252
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TYPE: PRT
ORGANISM: HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/60/160,209
CURRENT FILING DATE: 1999-10-19
NUMBER OF SEQ ID NOS: 4646
SOFTWARE: FastSEQ for Windows Version 4.0
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: ISOLATED HUMAN SECRETED PROTEINS,
HITLE OF INVENTION: UNCLEIC ACID MOLECULES ENCODING HUMAN SECRETED PROTEINS AND
HITLE OF INVENTION: USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: CL000113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: ISOLATE:
TITLE OF INVENTION: ACID MO:
TITLE OF INVENTION: THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PPLICANT: BONAZZI, VIVIEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: BONAZZI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
                                                                                                                                                                                                                                                                                                                 31 LLQRPGLQLYFS 42
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                                        374
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          VIVIEN
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                                                                                                                                                                                                                                                                                                                                                                                                        Score 61; DB 23;
Pred. No. 0.0018;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 61;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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. 0.0017;
                                                                                                                                                                                                                                                                                                                                                                                                                              DB 23;
                                                                                                                                                                                                                                                                                                                                                                                                                        Length 102;
                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 96;
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RESULT 6
US-08-332-562-94
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PCT-US00-05882-1128
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Best Local Similarity
Matches 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CT-US00-05882-1128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Ver. 2.0 SEQ ID NO 1128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 1128, Application PC/TUS0005882
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: PCT/US00/05882
CURRENT FILING DATE: 2000-03-08
EARLIER APPLICATION NUMBER: 60/124,270
EARLIER FILING DATE: 1999-03-12
NUMBER OF SEO ID NOS: 1694
                                                                                                                                                                                                                                                                                                    ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Craig Rosen, APPLICANT: Steve Ruben
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Human Cancer Associated Gene Sequences and Polypeptides
                        ZIP: 20007-5109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: SITE
LOCATION: (349)
OTHER INFORMATION: Xaa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE: PA106PCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LOCATION: (208)
OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
NAME/KEY: SITE
                                                                                            STREET: 5000
CITY: Washington
                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ENGTH: 399
                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                           APPLICANT:
                                                                                                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                              242 LLQRPGLQLYFS 253
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                             COUNTRY:
                                                                                                                                            ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               217 LLORPGLQLYFS 228
                                                                                                                                                                                                                                                                                                                                                                                                                  1 LLQRPGLQLYFS 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 LLQRPGLQLYFS 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12; Conservative
                                                                                                                                                                                           INVENTION:
                                                                                                                                                                                                                                                                                                                  Application US/08332562
                                                                                                                        3: Foley & Lardner
                                                                             USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                  HULETT, MAIK D.

VENTION: FC RECEPTOR FOR IMMUNOGLOBULIN EQUENCES: 130
                                                                                                                                                                                                                                                 MCKENZIE, Ian F.C.
HOGARTH, Mark P.
HIBBS, Margaret L.
                                                                                                                                                                                                                     BONADONNA, Lisa
                                                                                                                                                                                                                                       Bernadette M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               equals any of the naturally occurring
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 61;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 61; DB 21;
Pred. No. 0.0069;
                                                                                                                          Suite 500
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Query Match
Best Local Similarity
Matches 11; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NERAL INFORMATION
                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPETATING SYSTEM: PC-DOS/MS-DOS
                                                                            APPLICATION NUMBER: US 08/332,56
FILING DATE: 31-0CT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: AU PM8232/94
FILING DATE: 16-SEP-1994
                                                             RIOR APPLICATION DATA:
                                                                                                                                                        RIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Rel
                                                                                                                                                                                                                      URRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: (202)672-5300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RIOR APPLICATION DATA:
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                                              APPLICATION NUMBER:
                                                                                                                                                                         CLASSIFICATION: 435
                                                                                                                                                                                      APPLICATION NUMBER: US/08/462,973 FILING DATE: 05-JUN-1995
                                                                                                                                                                                                                                        SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                         ZIP: 20007-5109
                                                                                                                                                                                                                                                                                                                                                    STATE: D.C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 31-OC
                                                                                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ENGTH:
                                                                                                                                                                                                                                                                                                                                                                   Washington
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Application US/08462973
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3000 K Street, N.W.,
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                                                                                                                                                                                                                                                                                                                                       USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HOGARTH, Mark P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BONADONNA, Lisa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MCKENZIE, Ian F.C.
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27-MAY-1992
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                                 27-MAY-1992
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                                                                                                                                           US 08/332,562
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                                            US 07/896,457
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WO PCT/AU87/00159
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Pred. No. 0.013;
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TELEX: 904136
NFORMATION FOR SEQ ID NO:
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SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,
                                                                                                                                     FILING DATE: 25-OCT-:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
FILING DATE: 25-OCT--
                                                                                                                                                                                                                PRIOR APPLICATION DATA:
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                                                           REFERENCE/DOCKET NUMBER: 17
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29 LLQRPGLQLHFS 40
                            TELEFAX:
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                                                                                                                                                                                                                                             APPLICATION NUMBER: U. FILING DATE: 19920624
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                                            TELEPHONE:
                                                                                                                                                                                 APPLICATION NUMBER: PCT/1
FILING DATE: 25-OCT-1990.
                                                                                                                                                                                                                                  CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: AU PH6166 FILING DATE: 26-MAY-1986
                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                        20007-5109
                                                                                                                                                                                                                                                                                                                                                                                                   Washington, D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    : 91 amino acids amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                   3000 K Street, N.W., Suite 500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                : (202)672-5300
(202)672-5399
                            (202)672-5399
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HOGARTH, Phillip M.
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                                              (202)672-5300
                                                                                                                                     TOMBER: AU PJ 7045
25-OCT-1989
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                                                                           17227/111
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Pred. No.
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0.013;
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TYPE: AMINO ACID

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; MOLECULE TYPE: protein US-07-861-895-25
                                                                                                               CURRENT APPLICATION NUMBER: US/60/146,394
CURRENT FILING DATE: 1999-07-29
NUMBER OF SEQ ID NOS: 1866
SOUTWARE: FRASTSEQ for Windows Version 3.0
SEQ ID NO 1032
LENGTH: 373
                                                                                                                                                                                                                                                                            Sequence 1032, Application GENERAL INFORMATION:
APPLICANT: Bonazzi, Vivien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Ver. 2.0 SEQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                  60-146-394-1032
                                                                                                                                                                                                                       TITLE OF INVENTION: ISOLTITLE OF INVENTION: ACID
FEATURE: VARIANT
LOCATION: (1)...(373)
OTHER INFORMATION: Xaa =
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                                                                           ORGANISM: Drosophila
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PARLIER FILING DATE: 1998-02-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            URRENT APPLICATION NUMBER: US/09/245,764
URRENT FILING DATE: 1999-02-05
ARLIER APPLICATION NUMBER: 60/099,994
ARLIER FILING DATE: 1998-09-11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ENGTH: 261
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TLE OF INVENTION: THREE DATALLE OF INVENTION: AND USES THEREOF PROPRENCE: 4102-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          232 LLQRPGLQLHFS 243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LICANT: Powell, Maree S.
LICANT: McKenzie, Ian F.C.
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                                                                                                                                                                                                                                                                                                               Application US/60146394
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                                                                                                                                                                                                                                    ISOLATED DROSOPHILA PROTEINS, NUCLEIC ACID MOLECULES ENCODING DROSOPHILA PROTEINS, AND USES THERE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    THREE DIMENSIONAL STRUCTURES AND MODELS OF FC RECEPTORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             90.2%;
91.7%;
     Any Amino Acid
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Pred. No. 0
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NAME/KEY: VARIANT
COCATION: (1)...(611)
THER INFORMATION: Xaa = Any Amino Acid
US-60-146-055-774
                                                                                                                                                                                                                                                                                                                                                                                                                      US-60-146-055-774; Sequence 774; A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
Thes 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; ORGANISM: Klebsiella pneumoniae US-09-489-039A-13963
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             B
                                                 Query Match
Best Local S
                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 938
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 774
LENGTH: 611
                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS:
SEQ ID NO 13963
LENGTH: 535
TYPE: PRT
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                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/60/146,055
CURRENT FILING DATE: 1999-07-28
                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: ISOLATED SECRETED PROTEINS, TITLE OF INVENTION: MOLECULES ENCODING SECRETED
                                                                                                                                                                                                                                                                                                                                    FILE REFERENCE: CL000067
                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Bonazzi, Vivien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT FILING DATE: 2000-01-27
PRIOR APPLICATION NUMBER: US 60/117,747
PRIOR FILING DATE: 1999-01-29
NUMBER OF SEQ ID NOS: 14342
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/489,039A
CURRENT FILING DATE: 2000-01-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Gary Bre
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                TYPE: PRT
ORGANISM: Drosophila
                                                                                                                                                                                      FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       116 LLQRYGMQLY 125
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3 QRPGLQLYF 11
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                                                 Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gary Breton et. al
                                                                                                                                                                                                                                                                                                                                                                                                                     Application US/60146055
                                 Conservative
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                                                 63.9%; Score 39; 77.8%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        63.9%;
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7.7.8%;
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: Pred. No. 77;
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 39; DB 18;
Pred. No. 1.1e+02
                                 Mismatches
                                                 DB 23;
1.3e+02;
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77;
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                                                                                                                                                                                                                                                                                                                                                                  NUCLEIC ACID
                                                               Length 611;
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                                 Indels
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                               Gaps
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414 ERPDLQLYF 422

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TITLE OF INVENTION: ISOLATED G-PROTEIN COUPLED RECEPTORS,
TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING GPCR PROTEINS AND USES
TITLE OF INVENTION: THEREOF
FILE REFERENCE: CLOO0173
CURRENT APPLICATION NUMBER: US/60/173,464
CURRENT ILING DATE: 1999-12-29
NUMBER OF SEQ ID NOS: 30269
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 405
LENGTH: 612
TYPE: PRT
ORGANISM: Drosophila
US-60-173-464-405
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; Sequence 405; Application US/60173464
; GENERAL INFORMATION:
; APPLICANT: Li, Peter W.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; ORGANISM: Drosophila
US-60-167-217-452
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-60-167-217-452; Sequence 452, Application US/60167217
                                                                                                                                                                                                                                                                                                 RESULT 15
US-60-191-637-477
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 77.1
Matches 7; Conservative
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Best Local Similarity 77.9
Matches 7; Conservative
                                   Sequence 477, Application US/60191637
GENERAL INFORMATION:
APPLICANT: Venter, J. Craig
TITLE OF INVENTION: NUCLEIC ACID DETECTION KITS COMPRISING
TITLE OF INVENTION: GENE SEQUENCES EXPRESSED FROM THE DROSOPHILA GENOME,
TITLE OF INVENTION: USES THEREOF
FILE REFERENCE: CL000392
CURRENT APPLICATION NUMBER: US/60/191,637
CURRENT APPLICATION NUMBER: US/60/191,637
CURRENT FILING DATE: 2000-03-23
NUMBER OF SEQ ID NOS: 42660
SOFTMARE: FRANCESED for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/60/167,217 CURRENT FILING DATE: 1999-11-24 NUMBER OF SEQ ID NOS: 23195 SOFTWARE: FastSEQ for Windows Version 4.0 SEQ:ID NO 452 SEQ:ENGTH: 612 TYPE: PRT
LENGTH: 612
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REPLICANT: L1, Peter W. D.
RITLE OF INVENTION: ISOLATED DROSOPHILA PROTEINS, NUCLEIC
RITLE OF INVENTION: ACID MOLEUCLES ENCODING DROSOPHILA PROTEINS, AND USES
RITLE OF INVENTION: THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ILE REFERENCE:
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415 ERPDLQLYF 423
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77.88;
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ORGANISM: DROSOPHILA
US-60-191-637-477

QS-60-191-637-477

Query Match
Best Local Similarity 77.8%; Pred. No. 1.3e+02;
Matches 7; Conservative 1; Mismatches 1; Indels 0; Gaps

QY 3 QRPGLQLYF 11

QY 3 QRPGLQLYF 11
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Search completed: June 4, 2001, 12:20:02 Job time: 515 sec

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415 ERPDLOLYF 423

OM protein -

GenCore version 4.5 Copyright (c) 1993 - 2000 Comp

Compugen Ltd

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Result
No.
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Maximum DB
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Maximum Match 100%
Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Total number of hits satisfying chosen parameters:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Perfect score:
       Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                    Score
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seq.length: 2000000000
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61
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/cgn2_6/ptodata/1/paa/US06_NEW_COMB.pep:*
/cgn2_6/ptodata/1/paa/US07_NEW_COMB.pep:*
/cgn2_6/ptodata/1/paa/US08_NEW_COMB.pep:*
/cgn2_6/ptodata/1/paa/US09_NEW_COMB.pep:*
/cgn2_6/ptodata/1/paa/US09_NEW_COMB.pep:*
US-09-828-498-2

US-09-739-449-11020

US-09-724-475-3

US-09-769-066-21

US-09-79-49-11176

US-09-270-849B-19161

US-09-270-849B-189872

US-09-685-330-8

US-09-739-449-12557

PCT-US01-11988-1481

PCT-US01-11988-1482

US-09-685-330-6

US-09-685-330-6

US-09-685-330-6

US-09-833-245-1481

US-09-833-245-1481

US-09-833-245-1481
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US-09-270-849B-188156
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US-09-311-62
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9-449-11862
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1-626B-8
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12557, A
1481, Ap
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1864, Ap
1864, Ap
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11176, A
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186298,
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45	44	43	42	41	40	39	38	37	36	35	34	<u>3</u> 3	32	31	3 0	29	28
31	31	31	31	31	31	31	31	31	31	<u>3</u> 1	31	31	31	31	31	32	32
50.8	50.8	50.8	50.8	50.8	50.8	50.8	50.8	50.8	50.8	50.8	50.8	50.8	50.8	50.8	50.8	52.5	52.5
479	389	321	315	290	204	194	154	131	131	124	112	112	112	112	42	2356	387
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US-09-270-849B-193923	US-09-739-449-8238	US-09-270-849B-180893	US-60-248-505-1125	US-09-270-849B-182901	US-09-811-284-195	US-09-308-823A-197	PCT-US00-35017A-988	US-09-647-468-164	US-09-647-468-163	US-09-769-066-22	US-09-262-724-6	US-09-647-468-150	US-09-647-468-149	US-09-772-120-6	US-09-601-600-180	US-60-285-697-38	PCT-US00-35017A-1125
Sequence 193923,	Sequence 8238, Ap	Sequence 180893,	Sequence 1125, Ap	Sequence 182901,	Sequence 195, App	Sequence 197, App	•	•	•	Sequence 22, Appl	Sequence 6, Appli	Sequence 150, App	Sequence 149, App	Sequence 6, Appli	Sequence 180, App	Sequence 38, Appl	Sequence 1125, Ap

ALIGNMENTS

JS-09-284-107-23

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; OTHER INFORMATION: Oligopeptides representing the extracellular; OTHER INFORMATION: domain of CD64 US-09-284-107-23
                                                                                                                                                                                                                               US-09-270-849B-188156; Sequence 188156, App
CURRENT APPLICATION NUMBER: US/09/270;849B
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 195450
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 188156
LENGTH: 259
TYPE: PRT
                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Swimmer et al.
TITLE OF INVENTION: Insect
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 23, Application US/09284107 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 100
Best Local Similarity 100
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: PCT/NL97/00557
PRIOR FILING DATE: 1997-10-07
PRIOR APPLICATION NUMBER: EP 96202791.8
PRIOR FILING DATE: 1996-10-08
NUMBER OF SEQ ID NOS: 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Logtenberg, Ton
APPLICANT: de Kruif, Cornelis Adriaan John
TITLE OF INVENTION: METHODS AND MEANS FOR SELECTING PEPTIDES
TITLE OF INVENTION: AND PROTEINS HAVING SPECIFIC AFFINITY FOR A TARGET
FILE REFERENCE: 313632000600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT FILING DATE: 1999-10-25
                                                                                                                                             FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 12
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Pred. No.
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; ORGANISM: Homo sapiens
US-09-833-245-1866
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Best Local Similarity /2...
500 8; Conservative
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Best Local Similarity 55.6
Matches 10; Conservative
                                                                          SEQ ID NO 1866
                                                                        NUMBER OF SEQ ID NOS: 2267
SOFTWARE: PatentIn Ver. 2.1
EQ ID NO 1866
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Ver. SEQ ID NO 1866
                                                                                                                                                                                                                                                                                                                                                                         09-833-245-1866
                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/833,245
CURRENT FILING DATE: 2001-04-12
PRIOR APPLICATION NUMBER: 60/229, 358
PRIOR FILING DATE: 2000-04-12
PRIOR APPLICATION NUMBER: 60/256, 931
PRIOR FILING DATE: 2000-12-21
                                                                                                                                                 PRIOR APPLICATION NUMBER:
                                                      LENGTH:
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                                                                                                                                                                                                                                                                                                                    PPLICANT: Human Genome Sciences, Inc
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PRIOR APPLICATION NUMBER: 60/199, 384
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RIOR FILING DATE: 2000-04-12
                                                                                                                                                                                                                                                                                  ILE REFERENCE: PF546PCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           URRENT FILING DATE: 2001-01-12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ITLE OF INVENTION: Albumin Fusion Proteins ILE REFERENCE: PF546PCT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 3 LCQRPGLVLALPPQLSFS 20
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                                                                                                                                                 60/199; 384
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Pred. No. 1
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; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids US-09-833-245-1864
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                                                                                                                               SOFTWARE: PatentIn Ver.
SEQ ID NO 1864
LENGTH: 63
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Best Local Similarity 55.6%
Matches 10; Conservative
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Best Local Similarity
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Sequence 1864, A
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                                   NAME/KEY: SITE LOCATION: (37)
                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 60/256, 931 PRIOR FILING DATE: 2000-12-21
                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/199, 384 PRIOR FILING DATE: 2000-04-25
                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/833,245
CURRENT EILING DATE: 2001-04-12
PRIOR APPLICATION NUMBER: 60/229, 358
PRIOR FILLING DATE: 2000-04-12
                                                                                                                                                                                    NUMBER OF SEQ ID NOS:
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SOFTWARE: Patentin Ve
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PRIOR FILING DATE: 2000-12-21
PRIOR APPLICATION NUMBER: 60/199, 384
PRIOR FILING DATE: 2000-04-25
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                                                                                      ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                            TILE REFERENCE: PF546PCT
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LOCATION: (37)
DOTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
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                                                                                                               TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PPLICANT: Human Genome Sciences, Inc.
ITLE OF INVENTION: Albumin Fusion Proteins
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55.6%;
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Pred. No.
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Score 34;

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                                                                                                                                               ; OTHER INFORMATION: Description of Artificial Sequence: Synthetic
; OTHER INFORMATION: at any location having Xaa, Xaa means any amino acid
US-09-270-849B-186298
                                                                                                                                                                                                                                                                                                                                                                                                                                JS-09-270-849B-186298
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; ORGANISM: Sorangium cellulosum 
JS-09-311-626B-8
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Best Local Similarity
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LENCTH: 471
TYPE: PRT
                                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                              SEQ ID NO 186298
'LENGTH: 370
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Best Local Similarity
                                                                 Matches
                                                                                                                                                                                                                                                                             CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 195450
SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/270,849B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR FILING DATE: 1998-05-01
PRIOR APPLICATION NUMBER: 60/084,358
PRIOR FILING DATE: 1998-05-05
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                                                                                                                                                                                                             ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                  ILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                   PPLICANT: Swimmer et al
                                                                                                                                                                                                                                                                                                                                                             ITLE OF INVENTION: Insect genome survey devices
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PPLICANT: Schulein, Martin
PPLICANT: Outtrup, Helle
ITLE OF INVENTION: Novel Rhamnogalacturonan Hydrolases
                                                                                                                                                                                                 FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ILE REFERENCE: 5572.204-US
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243 RPNLXLYYS 251
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                             4 RPGLQLYFS 12
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                                                                               Score 33; DB 5;
Pred. No. 3.6e+02;
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Pred. No.
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                                                               Mismatches
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Best Local Similarity
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                                                                                                                                                Sequence 2, Application US/09828498 GENERAL INFORMATION:
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LENGTH: 393
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LENGTH: 391
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT FILING DATE: 2000-12-19
PRIOR APPLICATION NUMBER: US 09/514,000
PRIOR FILING DATE: 2000-02-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Hinkle, Gregory J.
APPLICANT: - Slater, Steven C.
TITLE OF INVENTION: Agrobacterium tumefaciens Genome Sequences and Uses Thereof
FILE REFERENCE: 38-10(15490)C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/739,449
CURRENT FILING DATE: 2000-12-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Slater, Steven C.
TITLE OF INVENTION: Agrobacterium tumefaciens Genome Sequences and Uses Thereof
FILE REFERENCE: 38-10(15490)C
                               APPLICANT: SCHMIDT, WARREN
APPLICANT: STAPLETON, JACK T.
TITLE OF INVENTION: FULL-LENGTH GB VIRUS C (HEPATITIS
TITLE OF INVENTION: ARE INFECTIOUS IN PRIMARY CD4 POS
                                                                                                             APPLICANT:
                                                                                                                                APPLICANT: XIANG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 13351
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/739,449
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Hinkle, Gregory J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
                     ILE REFERENCE:
URRENT APPLICATION NUMBER: US/09/828,498
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                                                                                                                                                                                                                                                               308 PGLQLY 313
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                                                                                                                                                                                                                                                                                                    5 PGLQLY 10
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                                                                                           WUNSCHMANN, SABI
SCHMIDT, WARREN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Application US/09739449
                       IOWA: 030US
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                                                                                                                                  JINHUA
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                        Score 33;
Pred. No.
                                                                                                                                                                                                                                                                                                                                          Mismatches
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                                                                                                                                                                                                                                                                                                                                                                      Length 393;
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                                                        RNA TRANSCRIPTS
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RESULT 13
US-09-724-475-3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Agrobacterium tumefaciens JS-09-739-449-11020
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; TYPE: PRT
; ORGANISM: Hepatitis G virus
US-09-828-498-2
                                                                                                                                                                                                                                                                                                                      Sequence 3, Application US/09724475 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity 50.1
Matches 7; Conservative
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SEQ ID NO 11020
LENGTH: 291
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Best Local Similarity
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CURRENT FILING DATE: 2000-12-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 09/514,000 PRIOR FILING DATE: 2000-02-23
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APPLICANT: Slater, Steven C.
TITLE OF INVENTION: Agrobacterium tumefaciens Genome Sequences and Uses Thereof
FILE REFERENCE: 38-10(15490)C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
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                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY DISK
                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: MORGAN & FINNEGAN, L.L.P.
                                                                                                                                                                                                                                                             TITLE OF INVENTION: Recombinant Proteins Of
                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                      APPLICANT: Tsarev,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER:
COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WORDPERFECT 5.1
                                                                                                                          CITY: NEW YORK
STATE: NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PatentIn Ver. 2.1
                                                                                          ZIP: 10154
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                                                                                                         USA
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Pred. No. 3
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Pred. No. 2
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Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21, Application US/09769066 GENERAL INFORMATION:
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (212) 7:
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        83 RPGLDLVFA 91
                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/542,634
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: Fabian, Gary R.
REGISTRATION NUMBER: 33,875
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4 RPGLQLYFS 12
                                REFERENCE/DOCKET NUMBER: 4600-0293.30
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 324-0880
                                                                                                                                                                                                        COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, V CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/769,066 FILING DATE: 24-Jan-2001 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                              ZIP: 94306
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Dehlinger & Associates
STREET: 350 Cambridge Ave., Suite
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: HEPATITIS E VIRUS ANTIGENS AND USES THEREFOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Fuerst, Thomas R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 758-4800
TELEFAX: (212) 751-6849
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA: US/09/724,475
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                    TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                              STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Palo Alto
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APPLICATION NUMBER: US08/316,765
FILING DATE: 03-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 28-Nov-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: 08/809,523
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Yarbough, Patrice O.
Zhang, Yifan
                    (415)
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                    324-0960
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Pred. No. 1.8e+02
                                                                                                                                                                                                                                                                                        Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                250
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Search completed: June 4, 2001, 12:21:29 Job time: 591 sec
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US-09-270-849B-191061
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORIGINAL SOURCE: SEQUENCE DESCRIPTION: SEQ ID NO: 21: US-09-769-066-21
                                                                                                                                                                                                                                                              ; OTHER INFORMATION: Description of Artificial Sequence: Synthetic US-09-270-849B-191061
                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/270,849B
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 195450
SOFTWARE: PATENTIN VET. 2.0
SEQ ID NO 191061
LENGTH: 141
TYPE: PRT
                                                                                                                                                                Query Match
Best Local Similarity 75.0
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 191061, Application US/09270849B
GENERAL INFORMATION:
APPLICANT: Swimmer et al.
TITLE OF INVENTION: Insect genome survey devices
FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 52.5%; Score 32; DB 5; Length 124; Best Local Similarity 66.7%; Pred. No. 1.8e+02; Matches 6; Conservative 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                    ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4 RPGLQLYFS 12
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83 RPGLDLVFA 91
                                                                                        1 LLQRPGLQ 8
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66 LIQRPGSQ 73
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LENGTH: 124 amino acids
TYPE: amino acid
STRANDEDNESS: Hepatitis E Virus (Burma Strain)
ORF-3
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75.0%;

    Mismatches

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GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.

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Result
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Maximum DB
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Maximum Match 100%
Listing first 45 summaries
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length: 2000000000
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                PCT-US00-05882-1128
US-08-332-562-94
US-08-462-973-94
US-07-861-895-25
                                                                                                    US-60-160-209-3193
US-60-160-209-4252
US-60-160-203-4414
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Sequence 1128, Ap
Sequence 94, Appl
Sequence 94, Appl
Sequence 25, Appl
Sequence 323, App
                                                                    Sequence 3193, Ap
Sequence 4252, Ap
Sequence 4414, Ap
Sequence 7, Appli
Sequence 14, Appl
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-60-185-362-7	-207-216-7	-60-185-361-5	-60-188-162-3	-60-177-646-2	-09-417-507-4	-60-170-430-2	-09-417-507-3	-60-170-430-2	-60-164-762-6	-60-196-190-1	-09-417-507-2	-09-417-507-3948	-60-155-811-8	-60-155-811-8	-60-155-811-5	-60-191-681-1802	-60-191-637-2286	-173-464-1	-60-191-681-1803	-60-191-681-1800	-60-191-637 [£] 2288	-60-191-637-2283	-60-173-464-1875	-60-173-464-1871	-60-167-217-2317	-60-167-217-2311	-60-167-217-2314	-09-270-767-4546	-60-167-217-2315	-US01-01324-2	-09-248-796-2	0-245-201-234	-09-417-5	38-3
722,	equence 747, Ap	equence 508,	uence 3505, A	equence	equence 42536,	equence 2462, A	equence	2622, A	equence 674, Ap	equence	equence	e 39489	equence 893,	equence	equence 561,	equence 1802	equence 22862,	Sequence 18737, A	equence 1803	equence 1800	equence 2288	equence 2283	equence 18758	18719,	equence 23174	equence 23115,	e 23147,	equence 4546	ce 23150	ce 2181,	equence 26957,	uence 234, Ap	equence 39462,	N

ALIGNMENTS

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; NUMBER OF SEQ ID NOS: 4
; SOFTWARE: FASTSEQ FOR W
; SEQ ID NO 3193
; LENGTH: 58
; TYPE: PRT
; ORGANISM: HUMAN
US-60-160-209-3193
US-60-160-209-4252; Sequence 4252, Application US/60160209; GENERAL INFORMATION:
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                                                            RESULT
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APPLICANT: BONAZZI, VIVIEN
APPLICANT: BONAZZI, VIVIEN
TITLE OF INVENTION: ISOLATED HUMAN KINASE PROTEINS, NUCLEIC
TITLE OF INVENTION: ACLD MOLECULES ENCODING HUMAN KINASE PROTIENS AND USES
TITLE OF INVENTION: THEREOF
FILE REFERENCE: CLOOO113
                                                                                                                                                                                            Query Match
Best Local Similarity
Matches: 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 3193,
                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/60/160,209
CURRENT FILING DATE: 1999-10-19
                                                                                                                1 FYMGSKTLRGRN 12
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25 FYMGSKTLRGRN 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Application US/60160209
                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                       for Windows Version 4.0
                                                                                                                                                                                                            100.0%;
                                                                                                                                                                                          Score 64; DB 23;
Pred. No. 7.5e-05;
Mismatches 0;
                                                                                                                                                                                                                                 Length 58;
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; ORGANISM: HUMAN
US-60-160-203-4414
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                                                                   APPLICANT: Hogarth, P. Mark
APPLICANT: Powell, Maree S.
APPLICANT: McKenzie, Ian F.C.
APPLICANT: Maxwell, Kelly F.
APPLICANT: Garrett, Thomas P.J.
APPLICANT: Epa, Vidana
                                                                                                                                                                                                                                                                                                                                     Ouery Match 100.0%;
Best Local Similarity 100.0%;
Matches 12; Conservative C
                                                                                                                                                                         GENERAL
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SEQ ID NO 4414
LENGTH: 102
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Best Local Similarity
Matches 12; Conserv
                          TITLE OF INVENTION: THREE DIMENSIONAL STRUCTURES AND MODELS OF FC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/60/160,203
CURRENT FILING DATE: 1999-10-19
NUMBER OF SEQ ID NOS: 6374
                     ILE REFERENCE: 4102-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 4252
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OTHER INFORMATION: X
-60-160-209-4252
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CURRENT FILING DATE: 1999-10-19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PPLICANT: BONAZZI, VIVIEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: BONAZZI, VIVIEN
TITLE OF INVENTION: ISOLATED
TITLE OF INVENTION: ACID MODERATED OF INVENTION: THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ILE REFERENCE: CL000116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: VARIANT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ENGTH:
                                                                                                                                                                                                                                                                       51 FYMGSKTLRGRN 62
                                                                                                                                                                                                                                                                                                         1 FYMGSKTLRGRN 12
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                                                                                                                                                                       INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INVENTION: USES THEREOF
                                                                                                                                                                                     Application US/09245764
CATION NUMBER: US/09/245,764
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ACID MOLECULES
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Pred. No. 0.00014;
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Pred. No. 0.
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                                          RECEPTORS
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                                                                                                                  SOFTWARE: PatentIn
SEQ ID NO 1128
LENGTH: 399
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; TYPE: PRT
; ORGANISM: HOMO sapiens
US-09-245-764-7
                                                                                                                                                                                                                                                                                                       Sequence 1128, Application GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                           CURRENT APPLICATION NUMBER: PCT/US00/05882
CURRENT FILING DATE: 2000-03-08
EARLIER APPLICATION NUMBER: 60/124/270
EARLIER FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 1694
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local
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Best Local Similarity
Matches 12; Conserv
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SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 7
                                                                                                                                                                                                                                                                     APPLICANT: Craig Rosen, APPLICANT: Steve Ruben
      LOCATION: (208)
OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
                                                                                    TYPE: PRT
ORGANISM: Homo sapiens
NAME/KEY: SITE
                                                          FEATURE:
NAME/KEY: SITE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/702,021
CURRENT FILING DATE: 2000-10-30
NUMBER OF SEQ ID NOS: 45
                                                                                                                                                                                                                                        TILE OF INVENTION: Human Cancer Associated TLE REFERENCE: PA106PCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: C. Frase
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  09-702-021-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EARLIER APPLICATION NUMBER: 60/099,994
EARLIER FILING DATE: 1998-09-11
EARLIER APPLICATION NUMBER: 60/073,972
EARLIER FILING DATE: 1998-02-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
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229 FYMGSKTLRGRN 24(
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les 12; Conser
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Pred. No.
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Pred. No. 0.00039;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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; LOCATION: (349) ; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids PCT-US00-05882-1128

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                                                                                  Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                           NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 542:
TELECOMPUNICATION INFORMATION:
TELEPHONE: (202)672-5300
TELEFAX: (202)672-5399
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            quence 94, Application US/08332562 ENERAL INFORMATION:
                                                                                                                                                                                                                  FORMATION FOR SEQ ID NO:
                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 91 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
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254 FYMGSKTLRGRN 265
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PPLICANT:
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1 FYMGSKTLRGRN 12
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41 FYVGSKILEYRN 52
                                                                        Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TLE OF INVENTION:
                                                                                                                                              TOPOLOGY: 11
                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US 07/896,457 FILING DATE: 27-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/0 FILING DATE: 31-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET:
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                                                                                                                                                                                                                                  904136
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3000 K Street, N.W., Suite 500
                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HIBBS, Margaret L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HULETT, Mark D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HOGARTH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MCKENZIE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BONADONNA, Lisa
                                                                                                                                              linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Foley & Lardner
                                                                                                                                                         single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bernadette M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                E, Ian F.C.
, Mark P.
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                                                                      62.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FC RECEPTOR FOR IMMUNOGLOBULIN
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                                                                                                                                                                                                                                                                                       54270/119/GRHA
                                                                        Score 40;
Pred. No.
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Pred. No.
                                                         Mismatches
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                                                                                     DB 7;
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                                                                                    Length 91;
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                                                         Indels
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RESULT 8 US-08-462-973-94

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US-07-861-895-25

GENERAL INFORMATION:

Application US/07861895

HOGARTH, Phillip M. HULETT, Mark D.

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Query Match
Best Local Similarity
Thes 8; Conserv
                                                                                                                                   US-08-462-973-94
                                                                                                                                                                                                                              INFORMATION FOR
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: 435
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 08/332,562
                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: Patentin Relaction DATA:
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                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATA:
PRIOR APPLICATION DATA:
AU PM8232/94
                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
41
                                                                                                                                                                LENGTH: 91 amino acids TYPE: amino acid STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: WO PCT/AU87/00159 FILING DATE: 29-MAY-1987
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Washington
                                                                                                                                                                                                                                                       TELEFAX:
                                                                                                                                                                                                                                                                         TELEPHONE:
                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                     NAME: BENT, Stephen A. REGISTRATION NUMBER: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
                            1 FYMGSKTLRGRN 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER:
FYVGSKILEYRN 52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20007-5109
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3000 K Street, N.W., Suite 500
                                                          Conservative
                                                                                                                                                                                                                                                         (202)672-5399
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HOGARTH, Mark P.
HIBBS, Margaret L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PatentIn Release #1.0, Version #1.30
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ENTION: FC RECEPTOR FOR IMMUNOGLOBULIN
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                                                                                                                                                  linear
                                                                                                                                                                                                                           SEQ ID NO:
                                                                                                                                                                                                                                                                         (202)672-5300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                05-JUN-1995
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26-MAY-1986
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                                                                       62.5%;
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                                                         pred. No. 5.4;
mismatches
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                                                                                        DB 8;
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IERINO, Francesco

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Best Local Similarity bo.,
Ahes 8; Conservative
                         FILE REFERENCE: 11000.1043U1
CURRENT APPLICATION NUMBER: US/09/634,238
CURRENT FILING DATE: 2000-08-08
                                                                                                                                                                                                                                                              SEQUENCE 323,
                                                               APPLICANT: Reid, Julian R.
APPLICANT: Coolbear, Timothy
TITLE OF INVENTION: Polynucleotides, materials incorporating
TITLE OF INVENTION: them and methods for using them.
                                                                                                                                                                                                                                                                                          09-634-238-323
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SEQUENCE CHARACTERISTICS:
LENGTH: 410 amino acid
TYPE: AMINO ACTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: protein -861-895-25
                                                                                                                                                                                                                                                                                                                                                244 FYVGSKILEYRN 255
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CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: AMINO ACID
TOPOLOGY: linear
LECHTE TO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: BENT, Stephen A. REGISTRATION NUMBER: 29 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                    1 FYMGSKTLRGRN 12
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FTLING DATE: 25-OCT-1990
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FILING DATE: 19920624
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                                                                                                                                                                                                                                                              INFORMATION:
                  SEQ ID NOS:
                                                                                                                                   Holland, Ross
O'Toole, Paul W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SSEE: Foley & Lardner
T: 3000 K Street, N.W.,
Washington, D.C.
                                                                                                                                                                  Christensson, Anna C.
                                                                                                                                                                                Lubbers, Mark
Dekker, James
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                                                                                                                                                                                                                                                                       Application US/09634238
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(202)672-5399
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                                                                                                                                                                                                                                              Matthew
for Windows Version
                                                                                                                                                                                                Mark W.
                                                                                                                                                                                                                                                                                                                                                                                                                        62.5%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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 4.0
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; ORGANISM: A.fumigatus US-09-417-507-39462
                                                                                                                                                     US-09-417-507-39462; Sequence 39462, Application US/09417507; GENERAL INFORMATION:
APPLICANT: KEITH G. WEINSTOCK ET AL.
                                                                                                                                                                                                                                                                    В
                              SEQ ID NO 39462
LENGTH: 197
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                            ; ORGANISM: Lactobacillus rhamnosus US-09-634-238-322
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Best Local Similarity
7; Conserv
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SEQ ID NO 322
LENGTH: 720
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                                                                                                                                                                                                                                                                                                                            Matches
                                                                                   CURRENT APPLICATION NUMBER: US/09/417,507 CURRENT FILING DATE: 1999-10-14
                                                                                                                     APPLICANT: KEITH G. WEINSTOCK ET AL.
TITLE OF INVENTION: FUMIGATUS FOR DIAGNOSTICS AND THERAPEUTICS
TITLE OF INVENTION: FUMIGATUS FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                        NUMBER OF SEQ ID NOS:
                                                                                                                  FILE REFERENCE: PATH99-10
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LENGTH: 268
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: CURRENT FILING DATE: 2000-0
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TITLE OF INVENTION:
FILE REFERENCE: 110
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                                                                                                                                                                                                                                                               407 YVGQKPLRSRN 417
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                                                                                                                                                                                                                                                                                           2 YMGSKTLRGRN 12
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                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                FastSEQ for Windows Version 4.0
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Bloksberg, Leonard,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Coolbear, Timothy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Reid, Julian R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              O'Toole, Paul W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Holland, Ross
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Dekker, James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Christensson, Anna
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Glenn, Matthew
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                                                                                                                                                                                                                                                                                                                                     59.4%;
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Pred. No.
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Pred. No. 43;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  materials incorporating for using them.
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                                                                                                                                                                                                                                                                                                                                           Length 720;
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                                                                                                                                                                                                                                                                                                                  Indels
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Query Match Best Local Similarity

57.8%; 45.5%;

Score 37; Pred. No.

DB 18;

Length 197;

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Query Match
Best Local Similarity
Best Local Similarity
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                                                                                                                                     PCT-US01-01324-2181
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US-09-248-796-26957
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JS-60-245-201-234
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 26957, Application US/09248796
GENERAL INFORMATION:
APPLICANT: Keith Weinstock et al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICAL
TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.132
                                                                                      Sequence 2181, Application PC/TUS0101324 GENERAL INFORMATION:
APPLICANT: Human Genome Sciences, Inc.
                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 7; Conserv
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LENGTH: 1845
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EQ ID NO 26957
LENGTH: 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/248,796
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TITLE OF INVENTION:
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                               PPLICANT: Human Genome Sciences, Inc., et al. ITLE OF INVENTION: Nucleic Acids, Proteins, a ILE REFERENCE: PCOC2PCT USO1/01324
cior application data removed - refer to PALM or file wrapper
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TLE OF INVENTION:
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89 FKIGSKTLEGQ 99
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DRUG-METABOLIZING PROTEINS,
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66.7%;
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Pred. No.
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PCT-US01-01324-2181
                                Best Local Similarity 87. Matches 7; Conservative
                                                               Query Match
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SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 2181
LENGTH: 153
                                                                                                                               OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
                                                                                                                                                  NAME/KEY: SITE LOCATION: (70)
                                                                                                                                                                                  OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Homo
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OTHER INFORMATION:
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4 GSKTLRGR 11
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Pred. No.
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                                                 DB 1;
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Search completed: June 4, 2001, 12:20:03 Job time: 516 sec

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Minimum DB
Maximum DB
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Perfect score:
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1    /cgn2_6/ptodata/1/paa/JCCT_NEW_COMB.pep:*

2    /cgn2_6/ptodata/1/paa/JUS06_NEW_COMB.pep:*

3    /cgn2_6/ptodata/1/paa/JUS07_NEW_COMB.pep:*

4    /cgn2_6/ptodata/1/paa/JUS08_NEW_COMB.pep:*

5    /cgn2_6/ptodata/1/paa/JUS09_NEW_COMB.pep:*
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Gapop 10.0 , Gapext 0.5
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US-09-525-361-29
PCT-US01-11986
              US-09-833-245-455

US-09-739-449-11123

US-09-603-124B-410

PCT-US00-01979A-71

US-09-493-795A-71

US-09-612-795-1

US-09-612-795-3

US-09-612-795-3

US-09-612-795-4
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US-09-493-795A-73

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US-09-739-449-12009

US-09-739-449-12004

US-09-739-449-12004

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PCT-US01-11988-458
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	Sequence 147, App	Sequence 63, Appl	Sequence 189576,	Sequence 1005, Ap	Sequence 1005, Ap	Sequence 2983, Ap	Sequence 1015, Ap	Sequence 190396,	Sequence 10658, A	Sequence 2, Appli	Sequence 11741, A	Sequence 11227, A	Sequence 139, App	Sequence 9493, Ap	Sequence 187832,	Sequence 194847,	Sequence 193304,	Sequence 3022, Ap
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ALIGNMENTS

US-09-284-107-24

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                                                                                                                                                                                                         Sequence 12, Application US/09459573
GENERAL INFORMATION:
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 24
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                                                                                                                                                                                                                                                         -09-459-573-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/284,107 CURRENT FILING DATE: 1999-10-25 PRIOR APPLICATION NUMBER: PCT/NL97/00557 PRIOR FILING DATE: 1997-10-07 PRIOR APPLICATION NUMBER: EP 96202791.8 PRIOR FILING DATE: 1996-10-08
                                                                                                                      APPLICANT: LIVSHITS, VITALIY
APPLICANT: ZAKATAEVA, NATALIA
APPLICANT: NAKANISHI, KAZUO
APPLICANT: VENIAMINOVICH, VLADIMIR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: de Kruif, Cornelis Adriaan John
TITLE OF INVENTION: METHODS AND MEANS FOR SI
TITLE OF INVENTION: AND PROTEINS HAVING SPI
FILE REFERENCE: 313632000600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Logtenberg, Ton
APPLICANT: de Kruif, Corn
                                                                                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT ORGANISM: Artificial Sequence
PPLICANT: TOKHMAKOVA, IRINA
TITLE OF INVENTION: METHOD FOR PRODUCING L-AMINO ACIDS
TEREFERENCE: 0010-1066-0
TURRENT APPLICATION NUMBER: US/09/459,573
TURRENT FILING DATE: 1999-12-13
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12; Conserv
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Pred. No.
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Query Match
Best Local Similarity
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ORGANISM: Conus radiatus CT-US00-01979A-73
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; Sequence 69, Application PC/TUS0001979A
; GENERAL INFORMATION:
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                                                                          SEQ ID NO 73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: PCT/US00/01979A
CURRENT FILING DATE: 2000-01-28
PRIOR APPLICATION NUMBER: US 60/118,381
PRIOR FILING DATE: 1999-01-29
NUMBER OF SEQ ID NOS: 404
SOFTWARE: PATENTIN VET: 2.0
SEQ ID NO 69
LENGTH: 61
                                                                                                                                                                                                                                               Sequence 73, Application PC/TUS0001979A
GENERAL INFORMATION:
APPLICANT: University of Utah Research Foundation
APPLICANT: Cognetix, Inc.
                                                                                        NUMBER OF SEQ ID NOS: 404
SOFTWARE: PatentIn Ver. 2.0
                                                                                                                            CURRENT FILING DATE: 2000-01-28
PRIOR APPLICATION NUMBER: US 60/118/381
PRIOR FILING DATE: 1999-01-29
                                                                                                                                                                     FILE REFERENCE: Alphas 2
CURRENT APPLICATION NUMBER: PCT/US00/01979A
CURRENT FILING DATE: 2000-01-28
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Best Local Similarity
Matches 8; Conserva
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TITLE OF INVENTION: Alpha-Conotoxin Peptides
FILE REFERENCE: Alphas 2
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SEQ ID NO 12
LENGTH: 212
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NUMBER OF SEQ ID NOS: 24
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                                                                                                                                                                                                                     PLICANT: Cognetix, Inc.
PLE OF INVENTION: Alpha-Conotoxin Peptides
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58.3%;
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Pred. No.
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Pred. No. 16;
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17 FTSGRSTFRGRN 28

1 FYMGSKTLRGRN 12

Matches

Query Match

Local

Similarity
7; Conserv

Conservative

54.7%;

Score 35; DB Pred. No. 11;

Length 61;

Mismatches

Indels

0;

Gaps

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US-09-493-795A-73
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                            NUMBER OF SEQ ID NOS: 404
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 73
LENGTH: 61
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
Thes 7; Conserv
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LENGTH: 61
TYPE: PRT
                                                                                                                                                                                                                                                                                                                  sequence /3,
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 54.7%;
Best Local Similarity . 58.3%;
Matches 7; Conservative
                                                                                                                                      APPLICANT: Jones, Robert M.
TITLE OF INVENTION: Alpha-Conotoxin Peptides
FILE REFERENCE: 2314-179. A
CURRENT APPLICATION NUMBER: US/09/493,795A
CURRENT FILING DATE: 2000-01-28
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                                                                                                           PRIOR APPLICATION NUMBER: US 60/118,381 PRIOR FILING DATE: 1999-01-29
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            ORGANISM: Conus radiatus
                                                                                                                                                                                                                                                                             NERAL INFORMATION:
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SOFTWARE: PatentIn Ver. 2.0
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PRIOR APPLICATION NUMBER: US 60/118,381
PRIOR FILING DATE: 1999-01-29
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CURRENT FILING DATE: 2000-01-28
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TITLE OF INVENTION: Alpha-Conotoxin Peptides
FILE REFERENCE: 2314-179.A
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McIntosh, J. Michael
                                                                                                                                                                                                                                                                                                                Application US/09493795A
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McIntosh, J. Michael
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Pred. No. 11;
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CT-US00-01979A-65

equence 65, Applica

Application PC/TUS0001979A

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SOFTWARE: PatentIn Ver. 2.
SEQ ID NO 65
LENGTH: 64
TYPE: PRT
ORGANISM: Conus textile
PCT-US00-01979A-65
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Best Local Similarity
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                                                                                            US-09-739-449-12089
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                                                               GENERAL INFORMATION:
                                                                            Sequence 12089, Application US/09739449
                                                                                                                                                                                                                                                   Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/493,795A
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NUMBER OF SEQ ID NOS: 404
                                               APPLICANT: Hinkle, Gregory J.
                                                                                                                                                                                                                                                                                                                                 LENGTH: 64
TYPE: PRT
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PRIOR APPLICATION NUMBER: US 60/118,381
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                                                                                                                                                                                                                      Local Similarity 58...
nes -7; Conservative
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                                                                                                                                                           FSSGRSTFRGRN 28
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              INVENTION: Agrobacterium tumefaciens Genome Sequences and Uses Thereof
                                                                                                                                                                                                                                                                                                                                                                                PatentIn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Jones, Robert M.
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                                  Slater,
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38-10(15490)C
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Query Match Best Local Similarity

51.6%; 54.5%;

Score 33; Pred. No.

DB 5; 24;

Length 55;

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; FEATURE: OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: at any location having Xaa, Xaa means any amir US-09-270-849B-194148
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Best Local Similarity
Thehes 6; Conserv
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US-09-270-849B-194148
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LENGTH: 377
                                                                                                                            SOFTWARE: Patentin Ver. SEQ ID NO 194148
                                                                                                                                                                                                                                                                                              Sequence 194148, Application US/09270849B
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Best Local Similarity
                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/270,849B
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 195450
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CURRENT FILING DATE: .2000-12-19
                                                                                                                                                                                                                                         APPLICANT: Swimmer et al TITLE OF INVENTION: Insec
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TITLE OF INVENTION: Agrobacterium tumefaciens Genome Sequences and Uses Thereof FILE REFERENCE: 38-10(15490)C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: US PRIOR FILING DATE: 2000-02-23
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                                                                                                                                                                                                                           FILE REFERENCE:
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TYPE: PRT
                                                                                           TYPE: PRT
                                                                        ORGANISM: Artificial Sequence
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Pred. No.
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Pred. No.
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Query Match
Best Local Similarity
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                                                          ORGANISM: Homo sapiens
CT-US01-11988-458
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US-09-823-356-16
           Query Match
                                                                                                                                      SEQ. ID NO 458
                                                                                                                                                  NUMBER OF SEQ ID NOS: 22
SOFTWARE: Patentin Ver.
                                                                                                                                                                                    PRIOR APPLICATION NUMBER: 60/256, 931
PRIOR FILING DATE: 2000-12-21
PRIOR APPLICATION NUMBER: 60/199, 384
PRIOR FILING DATE: 2000-04-25
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SOFTWARE: PERL Program
SEQ ID NO 16
LENGTH: 275
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                                                                                                                 ENGTH:
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URRENT FILING DATE: 2001-01-12
                                                                                                                                                                                                                                                                                                                                                             NERAL INFORMATION:
PPLICANT: Human Genome Sciences, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               URRENT APPLICATION NUMBER: US/09/823,356
URRENT FILING DATE: 2001-03-30
PRIOR APPLICATION NUMBER: 09/039,307
PRIOR FILING DATE: 1998 March 13
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Baughn, Mariah R.
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       51.6%;
                                                                                                                                                                                                                                                                        60/229, 358
   Score 33;
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Pred. No. 1.3e+02;
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 .DB 1;
Length 282
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Best Local Similarity bo...
Conservative
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US-09-833-245-458
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO 458
LENGTH: 282
TYPE: PRT
ORGANISM: Homo sapiens
                                                       PRIOR APPLICATION NUMBER: US 09/440,676 PRIOR FILING DATE: 1999-11-16
                                    PRIOR APPLICATION NUMBER: US 09/440,677
                                                                                                      PRIOR FILING DATE: 1999-11-15
PRIOR APPLICATION NUMBER: US 09/520,478
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         -09-525-361-29
                                                                                                                                          PRIOR APPLICATION NUMBER: US 09/440,493
                                                                                                                                                               PRIOR APPLICATION NUMBER: US 09/440,370 PRIOR FILING DATE: 1999-11-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity 85.7
Matches 6; Conservative
                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US 09/439,878
                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: US 09/268,865
PRIOR FILING DATE: 1999-03-15
PRIOR APPLICATION NUMBER: US 09/450,810
PRIOR APPLICATION NUMBER: US 09/453,137
PRIOR APPLICATION NUMBER: US 09/453,137
PRIOR APPLICATION NUMBER: US 09/453,137
                                                                                                                                                                                                                                                                                                                                          URRENT APPLICATION NUMBER: US/09/525,361
URRENT FILING DATE: 2000-03-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR FILING DATE: 2000-12-21
PRIOR APPLICATION NUMBER: 60/199, 384
PRIOR FILING DATE: 2000-04-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 2267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 60/229, 358
PRIOR FILING DATE: 2000-04-12
PRIOR APPLICATION NUMBER: 60/256, 931
                                                                                                                                                                                                                                                                                                                                                                                           ITLE OF INVENTION: NOVEL METHODS OF DIAGNOSING AND TITLE OF INVENTION: COMPOSITIONS, AND METHODS OF SCITLE OF INVENTION: CANCER MODULATORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                          PPLICANT: MACK, DAVID
PPLICANT: GISH, KURT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/833,245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Human Genome Sciences, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  equence 458, Applic
ENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        |uence 29, Application US/09525361
                                                                                                                                                                                                                                                                                                                                                                            LE REFERENCE: A-67860-3/DJB/JJD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ILE REFERENCE: PF546PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 46 YLGSKTL 52
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                      1999-11-16
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85.7%;
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1; Mismatches 0;
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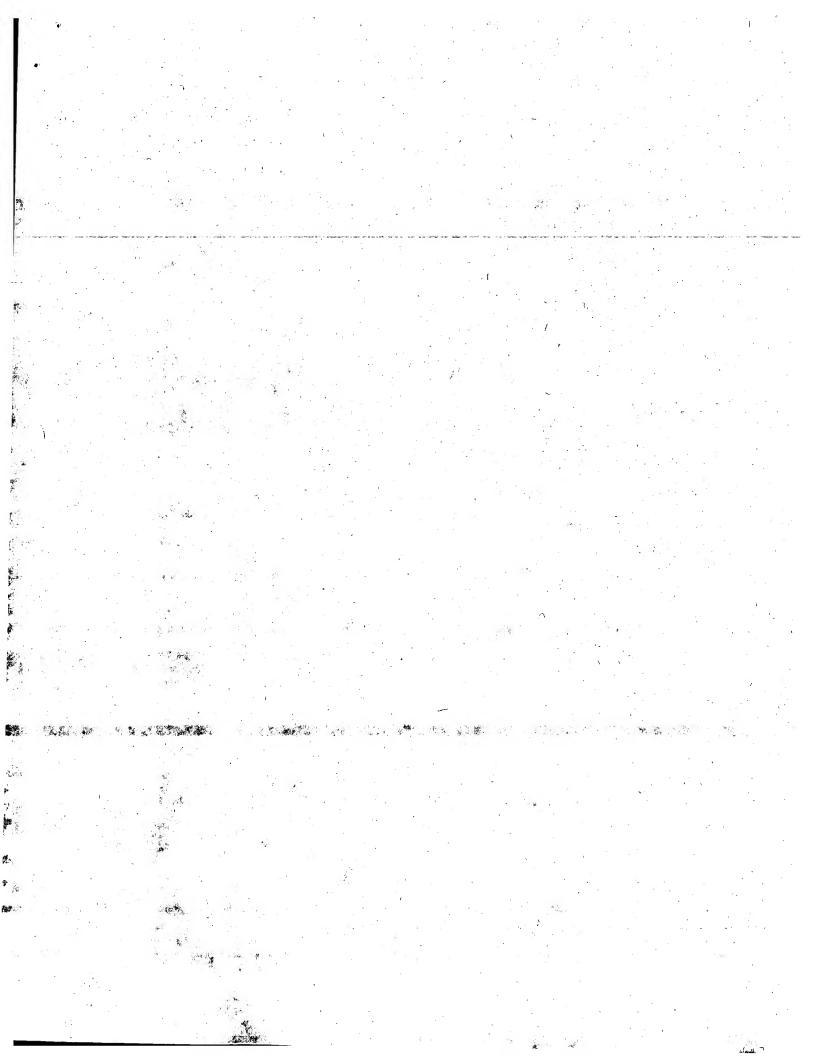
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: SEQ ID NO 29
; LENGTH: 291
; TYPE: PRT
; ORGANISW: Homo sapiens
US-09-525-361-29

Query Match
Best Local Similarity 85.7%; Pred. No. 1.4e+02;
Matches 6; Conservative 1; Mismatches 0; Indels 0; Gaps
Oy 2 YMGSKTL 8
i:||i||
Db 55 YLGSKTL 61

Search completed: June 4, 2001, 12:21:29

Search completed: June 4, 2001, 12:21:29
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GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.

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Result
No.
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Maximum DB
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Run on:
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                                                                                                                                                                                                                                                                                                           Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      seq length: 0 seq length: 2000000000
                                                                                                                                                                                                                     Match
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Gapop 10.0 , Gapext 0.5
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57
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                                                                                                                                                                                                                     Length
                                                           PCT-US00-05882-1128
US-60-173-468-1209
                                                                                                                                      US-60-160-209-3193
US-60-160-209-4252
US-60-160-203-4414
                                                                                                                                                                                                                                                                            SUMMARIES
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                                                                                                                                                                                                                                                                                                                                                                                                             _COMB.pep:
                                Sequence 3193, Ap
Sequence 4252, Ap
Sequence 4214, Ap
Sequence 7, Appli
Sequence 14, Appl
Sequence 1128, Ap
Sequence 1129, Ap
Sequence 621, App
                                                                                                                                                                                                                 Description
Sequence 94, Appl
Sequence 94, Appl
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-15135-14	-252-991A-	52-527	-US00-15135-15	35-15	-60-185-3	-08-993-(-08-824-1	-761-3		÷	-60-162-2	-09-248-7	۲	-09-417-5	-09-107-5	-60-191-681-	91-637-	-60-173-464-	-60-167-217-	-09-417-507-	-60-191-681-25	0-191-637-3	-60-173-464-25	-09-231-899-	-09-718-852-	-09-718-815-	-692-	S-09-25	07-861-	-09-718-8	-09-718-8	-09-718-692-4	-09-489-039A	16-6
Sequence 149, App	e 2167	e 5274	ئے	152,	615, 4	5167,	678, A	e 678,	σ.	e 6400, A		1490		e 37264,	Sequence 4715, Ap	6123,	e 7854,	6355,	Sequence 7951, Ap	equenc	Sequence 25591, A	Sequence 32284, A	Sequence 25166, A	Sequence 9, Appli	Sequence 2, Appli	Sequence 2, Appli			N			4, Ap	8221, A	•

ALIGNMENTS

US-60-160-209-4252 ; Sequence 4252, Application US/60160209 ; GENERAL INFORMATION: В Qy NUMBER OF SEQ ID NOS: 4646
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 3193
LENGTH: 58
TYPE: PRT
ORGANISM: HUMAN
US-60-160-209-3193 US-60-160-209-3193; Sequence 3193, Application US/60160209 RESULT Query Match
Best Local Similarity
Matches 12; Conser GENERAL INFORMATION: CURRENT APPLICATION NUMBER: US/60/160,209
CURRENT FILING DATE: 1999-10-19 APPLICANT: BONAZZI, VIVIEN
TITLE OF INVENTION: ISOLATED HUMAN KINASE PROTEINS, NUCLEIC
TITLE OF INVENTION: ACID. MOLECULES ENCODING HUMAN KINASE PROTIENS AND USES
TITLE OF INVENTION: THEREOF
FILE REFERENCE: CL000113 37 1 TSSEYQILTARR 12 TSSEYQILTARR 48 Conservative 100.0%; Score 57; DB 23; 100.0%; Pred. No. 0.00023; Mismatches Length 58; 0 Gaps

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    : NAME/KEY: VARIANT
: LOCATION: (1)...(96)
: OTHER INFORMATION: Xaa = Any Amino Acid
US-60-160-209-4252
                                                                                                                                                                                      Sequence 7, Application US/09245764 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                              Query Match 100.
Best Local Similarity 100.
Matches 12; Conservative
                       APPLICANT: Hogarth, P. Mark
APPLICANT: Howell, Maree S.
APPLICANT: McKenzie, Ian F.C.
APPLICANT: McKenzie, Ian F.C.
APPLICANT: Maxwell, Kelly F.
APPLICANT: Garrett, Thomas P.J.
APPLICANT: Epa, Vidana
TITLE OF INVENTION: THREE DIMENSIONAL STRUCTURES AND MODELS OF FC RECEPTORS
FILE REFERENCE: 4102-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 6374

SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4414

LENGTH: 102
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Sequence 4414, Application US/60160203
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Best Local Similarity
CURRENT APPLICATION NUMBER: US/09/245,764
CURRENT FILING DATE: 1999-02-05
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LENGTH: 96
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: HUMAN
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ITLE OF INVENTION:
ITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/60/160,209
CURRENT FILING DATE: 1999-10-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: ISOLATED HUMAN KINASE PROTEINS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PPLICANT:
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                                                                                                                                                                                                                                                                                              63 TSSEYQILTARR 74
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NUCLEIC ACID MOLECULES ENCODING HUMAN SECRETED PROTEINS AND
USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                100.0%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                Mismatches
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KINASE PROTIENS AND USES
                                                                                                                                                                                                                                                                                                                                                                                             Length 102;
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; SOFTWARE: PatentIn Ver.; SEQ ID NO 7; LENGTH: 261; TYPE: PRT; ORGANISM: Homo sapiens US-09-245-764-7
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US-09-702-021-14
; Sequence 14, Applica
; GENERAL INFORMATION:
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                                                                                                                             CURRENT APPLICATION NUMBER: PCT/US00/05882
CURRENT FILING DATE: 2000-03-08
EARLIER APPLICATION NUMBER: 60/124,270
EARLIER FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 1694
SOFTWARE: PATENTIN VET. 2.0
SEQ ID NO 1128
LENGTH: 399
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                    Sequence 1128, Application PC/TUS0005882 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
                                                                                                                                                                                                                                                          APPLICANT: Craig Rosen,
APPLICANT: Steve Ruben
TITLE OF INVENTION: Human Cancer Associated Gene Sequences and Polypeptides
FILE REFERENCE: PA106PCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                     NAME/KEY: SITE
LOCATION: (208)
OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/702,021
CURRENT FILING DATE: 2000-10-30
NUMBER OF SEQ ID NOS: 45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: C: Fras.
TITLE OF INVENTION:
FILE REFERENCE: 78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EARLIER APPLICATION NUMBER: 60/099,994
EARLIER FILING DATE: 1998-09-11
EARLIER APPLICATION NUMBER: 60/073,972
EARLIER FILING DATE: 1998-02-06
NUMBER OF SEQ ID NOS: 15
                                                                                  FEATURE:
                                                                                               ORGANISM: Homo sapiens
                                                                                                                  TYPE: PRT
NAME/KEY: SITE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                       241 TSSEYQILTARR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          222 TSSEYQILTARR 233
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 TSSEYQILTARR 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 TSSEYQILTARR 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C. Fraser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Application US/09702021
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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ilarity 100.0%;
Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ON: FAIL MOLECULES AND USES THEREOF 7853-217
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Pred. No. 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 57;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
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; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
PCT-US00-05882-1128
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                                                                                                                                              ; ORGANISM: HUMAN
US-60-181-426-621
                                                                                                                                                                                                                                                                                                                                                                                                  US-60-181-426-621
                                                                                                                                                                                                                                                                                                                                                  Sequence 621, Application U
GENERAL INFORMATION:
APPLICANT: BONAZZI, Vivien
                                                                Matches
                                                                               Query Match
Best Local 9
                                                                                                                                                                                         SEQ ID NO 62:
LENGTH: 56
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SOFTWARE: FastSEQ for
SEQ ID NO 1209
LENGTH: 56
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-60-173-468-1209
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                                                                                                                                                                                                                         CURRENT FILING DATE: 2000-02-09
NUMBER OF SEQ ID NOS: 708
SOFTWARE: FastSEQ for Windows V
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/60/173,468
CURRENT FILING DATE: 1999-12-29
                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/60/181,426
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                     ITLE OF INVENTION: ISOLATED HUMAN TRANSPORTER PROTEINS, ITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING HUMAN TRANSPORTER PROTEINS, ITLE OF INVENTION: AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PPLICANT: Bonazzi,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TLE OF INVENTION:
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les 7; Conserv
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                               2 SSEYQILTAR 11
SSEYETLTAK 24
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                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                     Application US/60181426
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illarity 100.0%;
Conservative
                                                                Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ISOLATED HUMAN DRUG TARGET PROTEINS, NUCLEIC ACID MOLECULES ENCODING HUMAN DRUG TARGET PROTIENS
                                                                               63.2%;
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                                                                               Score 36;
Pred. No.
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Pred. No.
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                                                                                           Length 56;
                                                               Indels
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RESULT

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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 91 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REGISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 54
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/896,457
FILING DATE: 27-MX-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                           CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
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                                                         NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                               53 TSSEYHIARAER 64
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX:
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                            ADDRESSEE:
                                                                                                                                                                                                                                                                                                            1 TSSEYQILTARR 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: BENT,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Washington
Washington
                                                                           INVENTION: FC RECEPTOR FOR IMMUNOGLOBULIN
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                                                                                                                                                                                                   Application US/08462973
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                3000 K Street, N.W., Suite 500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3000 K Street, N.W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HOGARTH, Mark P.
HIBBS, Margaret L.
                                                                                                         BONADONNA, Lisa
                                                                                                                                      HIBBS, Margaret L.
                                                                                                                                                                     MCKENZIE,
                                                                                                                                                                                                                                                                                                                                          Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SCOTT, Bernadette M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MCKENZIE,
                                                                                           HULETT, Mark D.
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                                                                                                                                                                                                                                                                                                                                                                                                                                    linear
                            Foley & Lardner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (202)672-5300
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                                                                                                                          Bernadette M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FC RECEPTOR FOR IMMUNOGLOBULIN
                                                                                                                                                        Mark
                                                                                                                                                                                                                                                                                                                                                         63.2%;
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                                                                                                                                                                                                                                                                                                                                                         Score 36;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Suite 500
                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Version #1.30
                                                                                                                                                                                                                                                                                                                                                                      Length 91;
                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                          0;
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OMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS

PatentIn Release #1.0, Version #1.30

ZIP: 20007-5109

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JS-60-207-216-686
                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity bu...
Ahes 8; Conservative
                                                           CURRENT APPLICATION NUMBER: US/60/207,216
CURRENT FILING DATE: 2000-05-26
NUMBER OF SEQ ID NOS: 898
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 686
                                                                                                                                       APPLICANT: Beasley,
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
FILE REFERENCE: CLOC
ORGANISM: HUMAN
-60-207-216-686
                                                                                                                                                                                                                                      Sequence 686,
                             LENGTH: 15
TYPE: PRT
                                                                                                                                                                                                                     ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 91 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: 16
TELECOMMUNICATION INFORMATION: (202)672-5300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/AU87/00159
FILING DATE: 29-MAY-1987
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/896,457
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APPLICATION NUMBER: US 08/332,562
APPLICATION STATE: 31-OCT-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: BENT, Stephen A. REGISTRATION NUMBER: 29,768
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                                                                                                                                                                                                                                                                                                                                              1 TSSEYQILTARR 12
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                                                151
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                                                                                                                                                                                                      Beasley, Ellen
                                                                                                                                                                                                                                 Application US/60207216
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                                                                                                                                                                  ISOLATED HUMAN DRUG TARGET PROTEINS NUCLEIC ACID MOLECULES ENCODING HUM
                                                                                                                                                      AND USES THEREOF
                                                                          Windows Version 4.0
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                                                                                                                                                                                                                                                                                                                                                                                                        DB 8;
                                                                                                                                                                ENCODING HUMAN DRUG TARGET PROTEINS,
                                                                                                                                                                                                                                                                                                                                                                                                        Length 91;
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Query Match

63.2%;

Score 36; DB 23;

Length 151;

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RESULT 14
US-09-718-815-4
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                                                             GENERAL INFORMATION:
                                                                            Sequence 4,
                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 4, Application US/09718692
GENERAL INFORMATION: APPLICANT: Beraud, Christophe
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Best Local 9
                                           APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/718,692
CURRENT FILING DATE: 2000-11-22
NUMBER OF SEQ ID NOS: 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Freedman, Richard
TITLE OF INVENTION: Novel motor proteins and methods for
TITLE OF INVENTION: their use
                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE:
                                                                                                                                                                                                                                                                                                                                LENGTH: 36
                                                                                                                                                                                                                                                                                                                                                                                                                                            FILE REFERENCE:
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CURRENT FILING DATE: 2000-01-27
PRIOR APPLICATION NUMBER: US 60/117,747
PRIOR FILING DATE: 1999-01-29
NUMBER OF SEQ ID NOS: 14342
                                                                                                                                                                                                                                                                                                                 ORGANISM: Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Gary Breton et. al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID
TITLE OF INVENTION: PNEUMONIAE FOR DIAGNOSTICS A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
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TYPE: PRT
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254 SSEYETLTAK 263
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                INVENTION:
                                                                                                                                                                                                                                                                                                                                               366
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7; Conservative
                                                                         Application US/09718815
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                                           Beraud, Christophe
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, Richard
Novel motor proteins and methods for
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58.3%;
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Pred. No. 42;
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2; Mismatches
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Sequence 4, Application US/09718852

GENERAL INFORMATION:
APPLICANT: Beraud, Christophe
APPLICANT: Ereedman, Richard
TITLE OF INVENTION: Novel motor proteins and methods for
TITLE OF INVENTION: their use
FILE REFERENCE: 1052
CURRENT APPLICATION NUMBER: US/09/718,852.
CURRENT FILING DATE: 2000-11-22
NUMBER OF SQO ID NOS: 4
SOFTWARE: FRASTSEQ for Windows Version 4.0
SEQ ID NO 4
LENGTH: 366
TYPE: PAT
ORGANISM: Human
US-09-718-852-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FITLE OF INVENTION: their use
FILE REFERENCE: 1052
CURRENT APPLICATION NUMBER: US/09/718,815
CURRENT FILING DATE: 2000-11-22
NUMBER OF SEQ ID NOS: 4
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4
LENGTH: 366
TYPE: PRT
ORGANISM: Human
US-09-718-815-4
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US-09-718-852-4
                                                                       Query Match
Best Local Similarity 70.0
Matches 7; Conservative
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Best Local Similarity 70.0%;
Matches 7; Conservative
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254 SSEYETLTAK 263
2 SSEYQILTAR 11
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254 SSEYETLTAK 263
                                                                                            63.2%; Score 36; DB 21; 70.0%; Pred. No. 48;
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Maximum Match 100%
Listing first 45 summaries
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Match
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57
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US-09-270-849B-180859
US-09-270-849B-190000
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1872, Ap
12993, A
180859,
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1460, Ap
190000,
12004, A
8153, Ap
4, Appli
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763, App
11547, A
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1753, Ap
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Sequence 6, Appl	Sequence 4, Appl	Sequence 1566, A	Sequence 8964, A	Sequence 1035, A	Sequence 3003, A	Sequence 581, App	Sequence 3665, A	Sequence 1697, A	Sequence 11488,	Sequence 188275,	Sequence 8568, A	Sequence 991, App	Sequence 1236, Ap	Sequence 89, App.	Sequence 2, Appl:	Sequence 4, Appli	Sequence 82, App.
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ALIGNMENTS

US-09-284-107-25

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; FEATURE:
OTHER INFORMATION: Oligopeptides representing the extracellular; OTHER INFORMATION: domain of CD64
US-09-284-107-25
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GENERAL INFORMATION:
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LENGTH: 12
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Best Local
APPLICANT: Hyseq Inc
TITLE OF INVENTION: Novel Nucleic Acids and Polypeptides
FILE REFERENCE: 784PCT
CURRENT APPLICATION NUMBER: PCT/US00/35017A
CURRENT FILING DATE: 2000-12-22
PRIOR APPLICATION NUMBER: US09/488,725
PRIOR APPLICATION NUMBER: US09/488,725
PRIOR FILING DATE: 2000-01-21
PRIOR FILING DATE: 2000-04-25
PRIOR FILING DATE: 2000-04-25
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SOFTWARE: FastSEQ for Windows Version 4.0
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CURRENT FILING DATE: 1999-10-25
PRIOR APPLICATION NUMBER: PCT/NL97/00557
PRIOR FILING DATE: 1997-10-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: de Kruif, Cornelis Adriaan John TITLE OF INVENTION: METHODS AND MEANS FOR STITLE OF INVENTION: AND PROTEINS HAVING SPFILE REFERENCE: 313632000600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Logtenberg, Ton APPLICANT: de Kruif, Corn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity 100 nes 12; Conservative
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PRIOR APPLICATION NUMBER: 60/229, 358
PRIOR APPLICATION NUMBER: 60/229, 358
PRIOR FILING DATE: 2000-04-12
PRIOR FILING DATE: 2000-12-21
PRIOR FILING DATE: 2000-12-21
PRIOR APPLICATION NUMBER: 60/199, 384
PRIOR FILING DATE: 2000-04-25
NUMBER OF SEC.
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Best Local Similarity
5; Conserve
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Best Local Similarity 54...
Thes 6; Conservative
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; TYPE: PRT
; ORGANISM: Homo sapiens
PCT-US01-06769-11
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PCT-US00-35017A-1460
                                                                                                                                              SOFTWARE: Patentin Ver. 2.1 SEQ ID NO 762
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CURRENT FILING DATE: 2001-03-02
PRIOR APPLICATION NUMBER: 60/186,519
PRIOR FILING DATE: 2000-03-02
NUMBER OF SEQ ID NOS: 13
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 11
OTHER INFORMATION: Xaa equals -US01-11988-762
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                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 2267
                                                                                                                                                                                                                                                                                                                                                                                                              -US01-11988-762 equence 762, Application PC/TUS0111988
                                 NAME/KEY: SITE
LOCATION: (110)
                                                                                 TYPE: PRT
ORGANISM: Homo sapiens
                                                                          FEATURE:
                                                                                                                                                                                                                                                                                                      URRENT APPLICATION NUMBER: PCT/US01/11988
URRENT FILING DATE: 2001-01-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Mayo Medical Ventures
TITLE OF INVENTION: hB7-H2, A NOVEL CO-STIMULATORY MOLECULE
FILE REFERENCE: 07039-202WO1
                                                                                                                              ENGTH:
                                                                                                                                                                                                                                                                                                                                                           PPLICANT: Human Genome Sciences, Inc.
ITLE OF INVENTION: Albumin Fusion Proteins
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54.5%;
   any of the naturally occurring L-amino acids
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Pred. No. 70;
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76;
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; ORGANISM: Homo sapiens
PCT-US01-11988-763
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; LOCATION: (110)
; COTHER INFORMATION: Xea equals any of the naturally occurring L-amino acids US-09-833-245-762
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; Sequence 763, App
 Query Match 56.1%;
Best Local Similarity 45.5%;
Matches 5; Conservative
                                                                                                                            SOFTWARE: PatentIn Ver.
SEQ ID NO 763
LENGTH: 290
TYPE: PRT
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Best Local Similarity 43...
5; Conservative
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Best Local Similarity 45...
""" hes 5; Conservative
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PRIOR FILING DATE: 2000-04-25
NUMBER OF SEQ ID NOS: 2267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 762
                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: PCT/US01/11988
CURRENT FILING DATE: 2001-01-12
PRIOR APPLICATION NUMBER: 60/229, 358
PRIOR FILING DATE: 2000-04-12
PRIOR APPLICATION NUMBER: 60/256, 931
PRIOR FILING DATE: 2000-12-21
PRIOR FILING DATE: 2000-12-21
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Sequence 762, A
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NUMBER OF SEQ ID NOS: 2267
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PRIOR APPLICATION NUMBER: 60/256, 931
PRIOR FILING DATE: 2000-12-21
PRIOR APPLICATION NUMBER: 60/199, 384
                                                                                                                                                                                                                                                                                                                                                               ITLE OF INVENTION: Albumin Fusion Proteins
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CURRENT FILING DATE: 2001-04-12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Human Genome Sciences, Inc.
TITLE OF INVENTION: Albumin Fusion Proteins
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45.5%;
Score 32; DB 1;
Pred. No. 96;
5; Mismatches
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Pred. No. 80;
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Pred. No. 80;
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                              Length 290;
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US-09-833-245-763

168 TSSDHQVLSGK 178

1 TSSEYQILTAR 11

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US-60-248-823-102; Sequence 102, Application US/60248823; GENERAL INFORMATION:
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Best Local Similarity
"---hes 7; Conserv
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Best Local Similarity
Thes 5; Conserve
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US-09-833-245-763
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SEQ ID NO 11547
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SEQ ID NO 763
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-739-449-11547
                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: US 09/514,000 PRIOR FILING DATE: 2000-02-23
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CURRENT FILING DATE: 2000-12-19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Slater, Steven C. ITTLE OF INVENTION: Agrobacterium tumefaciens Genome Sequences and Uses Thereof TLE REFERENCE: 38-10(15490)C
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PRIOR FILING DATE: 2000-12-21
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TYPE: PRT
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URRENT FILING DATE: 2001-04-12
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                                                                                                                     179 TALEFTILTAAR 190
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45.5%;
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Pred. No. 1.2e+02;
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Pred. No.
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Best Local Similarity
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US-09-739-449-8297
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US-60-248-823-102
                                                                                                                                                                                                                                                                                                                              PCT-US01-11988-133
SOFTWARE: Pat
SEQ ID NO 133
LENGTH: 359
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT FILING DATE: 2000-12-19
PRIOR APPLICATION NUMBER: US 09/514,000
PRIOR FILING DATE: 2000-02-23
NUMBER OF SEQ ID NOS: 13351
SEQ ID NO 8297
LENCTH: 240
TYPE: PRT
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 102
LENGTH: 715
TYPE: PRT
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Best Local (
                                                          NUMBER OF SEQ ID NOS:
                                                                                                                PRIOR APPLICATION NUMBER: 60/229, 358
PRIOR FILING DATE: 2000-04-12
PRIOR APPLICATION NUMBER: 60/256, 931
PRIOR FILING DATE: 2000-12-21
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TITLE OF INVENTION: Agrobacterium tumefaciens Genome Sequences and Uses Thereof
FILE REFERENCE: 38-10(15490)C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Hinkle, Gregory J. APPLICANT: Slater, Steven C.
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                                                                            PRIOR FILING DATE: 2000-04-25
                                                                                                                                                                                                                                TITLE OF INVENTION: Albumin Fusion Proteins FILE REFERENCE: PF546PCT
                                                                                                                                                                                                                                                                      APPLICANT: Human Genome Sciences, Inc.
                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: PCT/US01/11988
                                                                                                                                                                                                                                                                                                                                                                                                           149 TOSEYEIL 156
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                                                                                                APPLICATION NUMBER: 60/199,
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                                          PatentIn
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75.08;
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Pred. No.
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ORGANISM: Homo sapiens PCT-US01-11988-133

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Query Match
Best Local Similarity
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Best Local Similarity
Thes 7; Conserve
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                                                                                  RIOR APPLICATION NUMBER: 09/654,936
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Homo sapiens -09-833-245-133
                                                                                                                                                                                      JURRENT APPLICATION NUMBER: PCT/US01/04098A JURRENT FILING DATE: 2001-02-05 PRIOR APPLICATION NUMBER: NOT YET ASSIGNED PRIOR FILING DATE: 2001-01-30 PRIOR APPLICATION NUMBER: .09/728,422 PRIOR FILING DATE: 2000-11-30
                                                                                                                                                                                                                                                                                            APPLICANT: Hyseq, Inc.
FITLE OF INVENTION: NOVE
FILE REFERENCE: 21272-0;
                                                                                                                     RIOR APPLICATION NUMBER: 09/663,561
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                                                                                                                                                      RIOR APPLICATION NUMBER: 09/693,325
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-09-833-245-133
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NUMBER OF SEQ ID NOS: 2267
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PRIOR APPLICATION NUMBER: 60/229, 358
PRIOR FILING DATE: 2000-04-12
PRIOR FILING DATE: 2000-12-21
PRIOR FILING DATE: 2000-12-21
PRIOR APPLICATION NUMBER: 60/256, 931
PRIOR FILING DATE: 2000-12-21
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TITLE OF INVENTION: Albumin Fusion Proteins
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APPLICATION NUMBER: 09/620,325
FILING DATE: 2000-07-19
APPLICATION NUMBER: 09/598,075
FILING DATE: 2000-06-20
APPLICATION NUMBER: 09/560,875
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2 SSEYQILTA 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                 Novel Nucleic Acids and Polypeptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             54.48;
77.88;
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77.88;
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Pred. No. 2e+02;
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Pred. No. 2e+02;
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PCT-US01-04098A-1753; Sequence 1753, Application PC/TUS0104098A
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; ORGANISM: Mus musculus
US-09-193-562D-34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ď
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Best Local Similarity burners
6; Conservative
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                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Homo sapiens PCT-US01-04098A-1378
                                                                                                                                                                                                                                                                             FILE REFERENCE: 21272-029
CURRENT APPLICATION NUMBER: PCT/US01/04098A
CURRENT FILING DATE: 2001-02-05
                                                                                                            PRIOR FILING DATE:
                                                                                                                                                        PRIOR APPLICATION NUMBER: 09/663,561
                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 09/728,422
PRIOR FILING DATE: 2000-11-30
                                                                                                                                                                                                                                                                                                                         APPLICANT: Hyseq, Inc.
TITLE OF INVENTION: Novel Nucleic Acids and Polypeptides
FILE REFERENCE: 21272-029
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO 3
                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: Not yet Assigned PRIOR FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Pauli, Benedicht U.
TITLE OF INVENTION: Nucleotide Sequences Encoding Mammalian Calcium TITLE OF INVENTION: Activated Chloride Channel-Adhesion Molecules FILE REFERENCE: 18617.0052
CURRENT APPLICATION NUMBER: US/09/193,562D
CURRENT FILING DATE: 1998-11-17
PRIOR APPLICATION NUMBER: US/60/065,922
PRIOR APPLICATION NUMBER: US/60/065,922
NUMBER OF SEQ ID NOS: 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 34, Application US/09193562D GENERAL INFORMATION:
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Best Local Similarity
Matches 5; Conserv
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SOFTWARE: Custom
SEQ ID NO 1378
LENGTH: 790
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 902
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                                                                                                    APPLICATION NUMBER: 09/654,936 FILING DATE: 2000-09-01
                                                                                                                                                                                          APPLICATION NUMBER: 09/693,325
APPLICATION NUMBER: 09/560,875 FILING DATE: 2000-04-27
                                             APPLICATION NUMBER: 09/598,075
                                                                    APPLICATION NUMBER: 09/620,325 FILING DATE: 2000-07-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 TSSEYQILTA 10
                                                                                                                                          FILING DATE: 2000-09-15
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                                    2000-06-20
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Pred. No. 5.6e+02;
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Pred. No. 4.8e+02
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Run on:
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June 4, 2001, 12:20:03; Search time 260.2 Seconds (without alignments) 7.418 Million cell updates/sec
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OM protein - protein search, using sw model
                                                                                 GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.
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Title: Perfect score: Sequence: Scoring table: Searched: BLOSUM62 Gapop 10.0 , Gapext 0.5 US-09-284-107-26 64 1 EDSGLYQCEAAT 12

1009251 seqs, 160854530 residues

Total number of hits satisfying chosen parameters: 1009251

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Database : Pending_Patents_AA_Main:

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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SUMMARIES

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50	50	50	51	51	57	57	57	57	57	Score
78.1	78.1	78.1	79.7	79.7	89.1	89.1	89.1	89.1	89.1	Query Match Length DB ID
93	93	.93	81	81	. 399	374	261	102	96	ength
23	23	18	23	23	ب	21	16	23	23	DB .
US-60-173-469-1887	US-60-173-463-626	US-09-474-434-1676	US-60-177-571-3365	US-60-170-373-3410	PCT~US00-05882-1128	US-09-702-021-14	US-09-245-764-7	US-60-160-203-4414	3 US-60-160-209-4252	ID
Sequence 1887, Ap	Sequence 626, App	Sequence 1676, Ap	Sequence 3365, Ap	Sequence 3410, Ap	Sequence 1128, Ap	Sequence 14, Appl	Sequence 7, Appli	Sequence 4414, Ap	Sequence 4252, Ap	Description

	45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30 .	29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12	11
	47	47	47	47	48	48	48	48	48	48	48	48	4.8	48	48	48	48	48	48	48	48	48	50	50	50	50	50	50	50	50	50	50	50	50	50
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	US-60-196-710-6936		-60-170-374	70-373	-09-143-308	US-08-687-727B-4	US-09-143-308-5	US-08-687-727B-5	US-08-618-953-3	US-08-258-022-3	US-08-618-953-4	US-08-258-022-4	US-09-283-503-8	US-09-283-503A-8	PCT-US97-20201-8	US-09-173-151A-32	-09-073-363B-	٦	US-08-971-635-20	-09-040-714-	US-08-951-829-14	US-60-160-209-3193	-60-207-359-	-60 - 190	-60-190-000-	-60-186-656-	-60-186-656-	-60-186-282-	-60-186-282-	-60-186-280-	-60-173-686-	-60-173-469-	ò	-60-186-282-885	US-60-173-686-1676
	Sequence 6936, Ap					4	5	<u>ب</u>	Sequence 3, Appli	3, 2	e 4,	Sequence 4, Appli	80	Sequence 8, Appli	8, 1	32,	41,	19,	20,		14,	3193,	150,	280,	279, 1	1327,	1326,	543,		495, 7	1311,	1481,	1311,	885, 7	Sequence 1676, Ap

ALIGNMENTS

Oy 1 EDSGLYQCEAAT 12	Query Match Best Local Similarity • 91. Matches 11; Conservative	; OTHER INFORMATION: Xaa = US-60-160-209-4252		; NAME/KEY: VARIANT	; FEATURE:	; ORGANISM: HUMAN	; TYPE: PRT	; LENGTH: 96	; SEQ ID NO 4252	; SOFTWARE: FastSEQ for Windows Version 4.0	; NUMBER OF SEQ ID NOS: 4646	; CURRENT FILING DATE: 1999-10-19	; CURRENT APPLICATION NUMBER: US/60/160,209	; FILE REFERENCE: CL000113	; TITLE OF INVENTION: THEREOF	; TITLE OF INVENTION:	; TITLE OF INVENTION:	; APPLICANT: BONAZZI,	; GENERAL INFORMATION:	; Sequence 4252, Application US/60160209	US-60-160-209-4252	RESULT 1
12 78	89.1%; 91.7%; ative	(aa ≐ Any	_	:					`.	Windows	4646	1999-10-	WUMBER: U)113	HEREOF	CID MOLE	ISOLATED HUMAN	VIVIEN		tion US/		•
	Score 57; DB 23 Pred. No. 0.015; 0; Mismatches	Any Amino Acid								Version 4		.19	JS/60/160,2			CULES ENCO			-	60160209		
	; DB 23; . 0.015; tches 1	Δ.								.0			09			DING HUMAN	SE PROTEIN	•				
	Length 96;															ACID MOLECULES ENCODING HUMAN KINASE PROTIENS AND USES	KINASE PROTEINS, NUCLEIC					
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	Gaps								1							AND						
	0:															USES				,		

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RESULT 4
US-09-702-021-14
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US-09-245-764-7
                                      Sequence 14, Application US/09702021
GENERAL INFORMATION:
APPLICANT: C. Fraser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       В
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TITLE OF INVENTION: FAIL MOLECULES AND USES THEREOFFILE REFERENCE: 7853-217
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 15 SOFTWARE: PatentIn Ver.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 4414, Applicant INFORMATION:
APPLICANT: BONAZZI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 6374

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 4414

LENGTH: 102
                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                               Q ID NO 7
LENGTH: 261
                                                                                                                                                                                                                                                                                                                                                                                                                                              ARLIER FILING DATE: 1998-02-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PPLICANT: Epa, Vidana
ITTLE OF INVENTION: THREE DIMENSIONAL STRUCTURES AND MODELS OF
FITLE OF INVENTION: AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Maxwell, Kelly F.
APPLICANT: Garrett, Thomas P.J.
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APPLICANT: McKenzie, Ian F.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ARLIER APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: HUMAN
60-160-203-4414
                                                                                                                                                               1 EDSGLYQCEAAT 12
|||||| ||||
| 234 EDSGLYWCEAAT 245
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/60/160, 203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: ISOLATED HUMAN SECRETED PROTEINS,
TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING HUMAN SECRETED PROTEINS AND
TITLE OF INVENTION: USES THEREOF
                                                                                                                                                                                                                                       Local Similarity
nes 11; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 60/099,994
FILING DATE: 1998-09-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/09/245,764 FILING DATE: 1999-02-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Application US/09245764
                                                                                                                                                                                                                                    Conservative
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91.7%;
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                                                                                                                                                                                                                                               Score 57; DB 16
Pred. No. 0.041;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 57; [
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                                                                                                                                                                                                                                                                 DB 16;
                                                                                                                                                                                                                                                             Length 261;
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; FEATURE:
; NAME/KEY: SITE
; LOCATION: (349)
; OTHER INFORMATION: X
PCT-US00-05882-1128
                                                                                                                                                                                                                                             B
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                                                                                                                                                                                                                                                                                          Query Match

Best Local Similarity 91.7

Matches 11; Conservative
                                                                                                                                                Sequence 3410, Appli
GENERAL INFORMATION:
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CURRENT APPLICATION NUMBER: US/60/170,373
CURRENT FILING DATE: 1999-12-13
NUMBER OF SEQ ID NOS: 4282
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; ORGANISM: HOMO US-09-702-021-14
                                                                               TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                    FILE REFERENCE: CL000149
                                                                                                                                   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 1128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Human Cancer Associated Gene Sequences and Polypeptides FILE REFERENCE: PA106PCT CURRENT APPLICATION NUMBER: PCT/US00/05882 CURRENT FILING DATE: 2000-03-08 EARLIER APPLICATION NUMBER: 60/124,270 EARLIER FILING DATE: 1999-03-12
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Best Local Similarity
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NAME/KEY: SITE
LOCATION: (208)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 39
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/0: CURRENT FILING DATE: 2000-10-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                            278
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                                                                                                                                                                                                                                                                    1 EDSGLYQCEAAT 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    253 EDSGLYWCEAAT 264
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 EDSGLYQCEAAT 12
                                                                                                                                                                                                                                     EDSGLYWCEAAT 289
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            399
                                                                                                                               Bonazzi,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FastSEQ for Windows Version 3.0
                                                                                                                                                          Application US/60170373
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                                                                                      ISOLATED HUMAN KINASE PROTEINS, NUCLEIC ACID MOLECULES ENCODING HUMAN KINASE PRO
                                                                                                                              Vivien
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91.78;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                             DB 1;
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0.059;
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                                                                                                                                                                                                                                                                                                                                                                                    occurring L-amino acids
                                                                                      PROTIENS AND USES
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; ORGANISM: Human US-60-170-373-3410

TYPE: PRT

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Best Local Similarity
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SOFTWARE: FastSEQ for w
SEQ ID NO 1676
SEQ ID NO 1676
LENGTH: 93
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                JS-09-474-434-1676
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JS-60-177-571-3365
                            Query Match
Best Local S
Matches 9
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CURRENT FILING DATE: 2000-01-27
NUMBER OF SEO ID NOS: 5082
SOFTWARE: FAStSEQ for Windows Version 4.0
SEQ ID NO 3365
LENGTH: 81
TYPE: PRT
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Best Local (
                                                                                                        -09-474-434-1676
                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/474,434
CURRENT FILING DATE: 1999-12-28
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                                                                                                                                                                                                                                                                                                                                                                                                 equence 1676, Application US/09474434
                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: ISOLATED HUMAN TITLE OF INVENTION: ACID MOLECULES ITLE OF INVENTION: THEREOF
                                                                                                                                                                                                                                                                                                                                                                                 ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ENERAL INFORMATION:
                                                                                                                  NAME/KEY: VARIANT
LOCATION: (1)...(93)
OTHER INFORMATION: Xaa = Any Amino Acid
                                                                                                                                                                                                                                                                                                         'ILE REFERENCE:
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                                                                                                                                                                    FEATURE:
                                                                                                                                                                                 ORGANISM: HUMAN
                                                                                                                                                                                                                                                                                                                                                                    PPLICANT: BONAZZI, VIVIEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PPLICANT: Bonazzi,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LE REFERENCE: CL0002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             56 EDAGEYQCEAST 67
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EDSGLYQCEA 10
                              Similarity
9; Conser
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9; Conser
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                              Conservative
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                                                                                                                                                                                                                                             Windows Version 4.0
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75.08;
                                            78.1%;
90.0%;
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75.08;
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Pred. No.
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                            Score 50; DB 1
Pred. No. 0.25;
0; Mismatches
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                                                          DB 18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 81;
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                                                         Length 93;
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                              Indels
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Query Match
Best Local Similarity
Thes 9; Conserve
                                                                            US-60-173-686-1676; Sequence 1676, Application US/60173686
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; ORGANISM: HUMAN
US-60-173-469-1887
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US-60-173-463-626
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                                                                                                                    RESULT
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COCATION: (1)...(93)
OTHER INFORMATION: Xaa =
US-60-173-463-626
                                                GENERAL INFORMATION:
APPLICANT: BONAZZI,
                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 2120
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1887
LENGTH: 93
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APPLICANT: BONAZZI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 702
SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 626
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TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/60/173,469
CURRENT FILING DATE: 1999-12-29
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TITLE OF INVENTION: ISOLATED HUMAN ION CHAN
TITLE OF INVENTION: NUCLEIC ACID MOLECULES
TITLE OF INVENTION: AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: ISOLATED HUMAN SECRETED PROTEINS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ITLE OF INVENTION:
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nes 9; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE: CL000185
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               ISOLATED HUMAN KINASE PROTEINS, NUCLEIC AND USES ACID MOLECULES ENCODING HUMAN KINASE PROTIENS AND USES
                                                  VIVIEN
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90.08;
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Pred. No. 0.
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                                                                                                                                                                                                                                                                   Length 93;
                                                                                                                                                                                                                                   Indels
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APPLICANT: BONAZZI, VIVIEN
TITLE OF INVENTION: ISOLATED HUMAN KINASE PR
TITLE OF INVENTION: ACID MOLECULES ENCODING
TILE OF INVENTION: THEREOF
FILE REFERENCE: CL000182
CURRENT APPLICATION.NUMBER: US/09/474,434
CURRENT FILING DATE: 1999-12-28
NUMBER OF SEO ID NOS: 1872
SOFTWARE: FastSEO for Windows Version 4.0
SEQ ID NO 1311
                                                                                                                                                                                      RESULT 13
US-09-474-434-1311
; Sequence 1311, Applic
; GENERAL INFORMATION:
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: OTHER INFORMATION: Xaa = Any Amino
US-60-186-282-885
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LOCATION: (1)...(93)
OTHER INFORMATION: X
US-60-173-686-1676
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Best Local Similarity 90.0
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Best Local S
Matches 9
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CURRENT FILING DATE: 2000-03-01
NUMBER OF SEQ ID NOS: 896
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 885
LENGTH: 93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           S-60-186-282-885
Sequence 885, Application US/60186282
GENERAL INFORMATION:
APPLICANT: BONAZZI, Vivien
TITLE OF INVENTION: ISOLATED HUMAN SECRETED PROTEINS,
TITLE OF INVENTION: USCLETIC ACID MOLECULES ENCODING HUMAN SECRETED PROTEINS, AND
TITLE OF INVENTION: USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: CL000182
CURRENT APPLICATION NUMBER: US/60/173,686
CURRENT FILING DATE: 1999-12-29
NUMBER OF SEQ ID NOS: 1872
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1676
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: HUMAN
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90.0%;
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Pred. No. 0.25;
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Pred. No. 0:25;
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                                                                                                                           KINASE PROTEINS, NUCLEIC ENCODING HUMAN KINASE PROTIENS AND USES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
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g 8

70 EDSGTYQCEA 79

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; LENGTH: 95
; TYPE: PRT
; ORGANISM: HUMAN
US-60-173-469-1481
               Query Match
Best Local Similarity
---hes 9; Conserv:
                                                                                                  ; ORGANISM: HUMAN US-60-173-686-1311
                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
Finance 9; Conserv
                                                                                                                                          NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ for
SEQ ID NO 1311
LENGTH: 95
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; ORGANISM: HUMAN
US-09-474-434-1311
                                                                                                                                                                                                                                                                                                          Sequence 1311, Application US/60173686 GENERAL INFORMATION:
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Best Local Similarity 90.0
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CURRENT FILING DATE: 1999-12-29
NUMBER OF SEQ ID NOS: 2120
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1481
                                                                                                                                                                                                                          TITLE OF INVENTION: ISOLATED HUMAN KINASE PROTEINS, NUCLEIC TITLE OF INVENTION: ACID MOLECULES ENCODING HUMAN KINASE PROTIENS AND USES FILE REFERENCE: CL000182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
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                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/60/173,686
CURRENT FILING DATE: 1999-12-29
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TITLE OF INVENTION: ISOLATED HUMI
TITLE OF INVENTION: USES THEREOF
                                                                                                                                    TYPE: PRT
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1 EDSGLYQCEA 10
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90.0%;
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Pred. No.
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                                           0.25;
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Search completed: June 4, 2001, 12:20:04 Job time: 517 sec

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1 PCT-US01-09226-47
                      US-09-819-136-2
PCT-US01-11988-2222
US-09-833-245-2222
US-09-814-950-4
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US-09-809-391-759
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Sequence 10, Appl
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Sequence 759, App
Sequence 759, App
Sequence 188150,
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Sequence 11, Appl
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	Sequence 3, Appli	Sequence 19, Appl	Sequence 17, Appl	Sequence 15, Appl	Sequence 8, Appli	Sequence 5, Appli.	Sequence 6, Appli	Sequence 20, Appl	Sequence 14, Appl	Sequence 13, Appl	Sequence 186367,	Sequence 186137,	Sequence 182954;	Sequence 190337,	Sequence 5, Appli	Sequence 3, Appli	Sequence 14, Appl	Sequence 7, Appli

ALIGNMENTS

OTHER INFORMATION: Oligopeptides representing the extracellular OTHER INFORMATION: domain of CD64 US-09-284-107-26 US-09-284-107-26 Sequence 26, Application US/09284107 GENERAL INFORMATION: APPLICANT: Logtenberg, Ton APPLICANT: de Kruif, Cornelis Adria NUMBER OF SEQ ID NOS: 29 SOFTWARE: PastSEQ for Windows Version 4.0 SEQ ID NO 26 LENGTH: 12 CURRENT APPLICATION NUMBER: US/09/284,107 CURRENT FILING DATE: 1999-10-25 PRIOR APPLICATION NUMBER: PCT/NL97/00557 PRIOR FILING DATE: 1997-10-07 PRIOR APPLICATION NUMBER: EP 96202791.8 -PRIOR FILING DATE: 1996-10-08 APPLICANT: de Kruif, Cornelis Adriaan John TITLE OF INVENTION: METHODS AND MEANS FOR SELECTING PEPTIDES TITLE OF INVENTION: AND PROTEINS HAVING SPECIFIC AFFINITY FOR A TARGET FILE REFERENCE: 313632000600 ORGANISM: Artificial Sequence FEATURE:

밁 Ş PCT-US01-10462-11 RESULT Query Match Best Local Similarity Matches 12; Conserv Sequence 11, Application PC/TUS0110462 GENERAL INFORMATION: APPLICANT: Liu, Chenghua
APPLICANT: Tang, Y. Tom
APPLICANT: Drmanac, Radoje T.
APPLICANT: Drmanac, Radoje T.
TITLE OF INVENTION: METHODS AND MATERIALS RELATING TO INSULIN-LIKE GROWTH FACTOR B
TITLE OF INVENTION: PROTEIN-LIKE POLYPEPTIDES AND POLYNUCLEOTIDES APPLICANT: APPLICANT: Yamaza URRENT APPLICATION NUMBER: PCT/US01/10462 1 EDSGLYQCEAAT 12 ||||||||||| 1 EDSGLYQCEAAT 12 Yamazaki, Vici Asundi, Vinod Conservative INC. 100.0%; Score 64; 100.0%; Pred. No. 0, Mismatches 1.6e-05; Length 12; Indels 0, Gaps 0,

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Query Match
Best Local Similarity
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                                                                                      GENERAL INFORMATION:
                                                                                                     Sequence 6, Application PC/TUS0110462
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LENGTH: 251
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SOFTWARE: PatentIn version 3.0
SEQ ID NO 11
LENGTH: 103
TYPE: PRT
                                   APPLICANT:
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                                                                     APPLICANT: HYSEQ, INC.
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: 09/540,217
PRIOR FILING DATE: 2000-03-31
NUMBER OF SEQ ID NOS: 14
                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT ORGANISM: homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      URRENT APPLICATION NUMBER: PCT/US01/10462
URRENT FILING DATE: 2001-03-30
PRIOR APPLICATION NUMBER: 09/784,748
PRIOR FILING DATE: 2001-02-14
PRIOR APPLICATION NUMBER: 09/649,167
PRIOR FILING DATE: 2000-08-23
PRIOR FILING DATE: 2000-08-23
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PPPLICANT: Drmanac, Radoje T.
PPPLICANT: Drmanac, Radoje T.
PRITICE OF INVENTION: METHODS AND MATERIALS RELATING TO INSULIN-LIKE GROWTH FACTOR BIND
PRITICE OF INVENTION: PROTEIN-LIKE POLYPEPTIDES AND POLYNUCLEOTIDES
PROTEIN-LIKE POLYPEPTIDES AND POLYNUCLEOTIDES
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PRIOR FILING DATE: 2000-08-23
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                                Yamazaki, Victoria
Asundi, Vinod
                Liu, Chenghua
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Pred. No. 0.46
1; Mismatches
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Pred. No.
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SEQ ID NO 759 LENGTH: 198

Prior application data removed - NUMBER OF SEQ ID NOS: 761

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SOFTWARE: PatentIn Ver.

CURRENT APPLICATION NUMBER: US/09/809,391
CURRENT FILING DATE: 2001-03-16

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US-09-809-391-759
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GENERAL INFORMATION:
APPLICANT: Ruben et al.
TITLE OF INVENTION: 186 Human Secreted proteins
FILE REFERENCE: PZ002P2.
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Best Local Similarity 100
Conservative
                                                                              Sequence 759,
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SEQ ID NO 760
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CURRENT FILING DATE: 2001-03-16
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PRIOR APPLICATION NUMBER: 09/784,748
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TITLE OF INVENTION: METHODS AND MATERIALS RELATING TO INSULIN-LIKE GROWTH FACTOR B
TITLE OF INVENTION: PROTEIN-LIKE POLYPEPTIDES AND POLYNUCLEOTIDES
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TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        186 Human Secreted proteins
                                                                                                                                                                                                                                                           71.9%; Score 46; 100.0%; Pred. No.
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Pred. No.

    consult PALM or file wrapper

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                                                                                                                                                                                                                                                             DB 5;
0.47;
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Query Match
Best Local Similarity
Conserv

Conservative

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71.9%;

DB 5;

Length 198;

0

Indels

Gaps

; ORGANISM: Homo sapiens US-09-809-391-759

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RESULT 9
US-09-142-956B-14
                                                                                                                                                                                                                              ; OTHER INFORMATION: Description of Artificial Sequence: Synthetic; OTHER INFORMATION: at any location having Xaa, Xaa means any amin US-09-270-849B-188150.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-09-270-849B-188150; Sequence 188150, Application US/09270849B; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                JS-09-809-391-478
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 100.
Ches 8; Conservative
                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 195450
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 188150
                                                                                                                                                                        Query Match
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SEQ ID NO 478
LENGTH: 234
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CURRENT FILING DATE: 2001-03-16
                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 09-809-391-478
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: 186 Human Secreted proteins FILE REFERENCE: PZ002P2
                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/270,849B
                                                                                                                                                                                                                                                                                                                                                                                                                                             TLE REFERENCE:
                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                      TYPE: PRT
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PPLICANT: Ruben et al
                                                                                                                                                                                                                                                                                   FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PPLICANT: Swimmer et al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            ITLE OF INVENTION: Insect genome survey devices
                                                                                                                                                                                                                                                                                                                                    ENGTH: 261
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         106 EDSGLYQC 113
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                                                                   2 DSGLYQCEAAT 12
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68 DSGIYECQVST 78
                                                                                                                                                          Local Similarity
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54.58;
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Pred. No. 0.65;
                                                                                                                                                                        Score 44;
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                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                          DB 5;
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Sequence 14, Application US/09815626 GENERAL INFORMATION:

APPLICANT: Glucksmann, Maria Alexandra
TITLE OF INVENTION: 33395, A NOVEL HUMAN LEUCINE-RITITLE OF INVENTION: FAMILY MEMBER AND USES THEREOF

HUMAN LEUCINE-RICH REPEAT

CURRENT APPLICATION NUMBER: US/09/815,626
CURRENT FILING DATE: 2001-03-23

URRENT FILING DATE: 2001-03-23
RIOR APPLICATION NUMBER: US 60/191,863
RIOR FILING DATE: 2000-03-24

'ILE REFERENCE: 10448-031001

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Best Local Similarity
Tropics 7; Conserve
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Best Local Similarity
Thes 7; Conserv:
RESULT 11
US-09-815-626-14
                                                                                                                                                                                                                      ; ORGANISM: Homo sapien 
US-09-232-773A-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GRGANISM: Homo sapiens US-09-142-956B-14
                                                                                                                                                                                                                                                                                     SOFTWARE: FastSEQ for
SEQ ID NO 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/232,773A CURRENT FILING DATE: 1999-07-15
                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Kendall, Richard L.
APPLICANT: Thomas, Kenneth A., Jr.
TITLE OF INVENTION: INHIBITOR OF VASCULAR ENDOTHELIAL CELL
TITLE OF INVENTION: GROWTH FACTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ULT 10
09-232-773A-15
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                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                           FILE REFERENCE: 18888DB
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TITLE OF INVENTION: VEGF-BINDING POLYPEPTIDES
FILE REFERENCE: 06501-021001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Niwa, Mikio
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TYPE: PRT
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                                                               730 EDEGLYTCOACS 741
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                                                                                                                                   Conservative
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                                                                                                                                                                                                                                                                                                       Windows Version 4.0
                                                                                                                                                     67.2%;
58.3%;
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Pred. No. 8;
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Pred. No. 7
                                                                                                                                     Mismatches
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Best Local Similarity
"" Researches 8; Conserve
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; OTHER INFORMATION: consensus sequence
US-09-815-626-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                              Sequence 13, Application PC/TUS0110462 GENERAL INFORMATION:
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CURRENT FILING DATE: 2001-03-30
PRIOR APPLICATION NUMBER: US 60/193,919
PRIOR FILING DATE: 2000-03-31
NUMBER OF SEQ ID NOS: 10
SOFTMARE: FastSEQ for Windows Version 4.0
SEQ ID NO 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 9, Application US/09822687
GENERAL INFORMATION:
APPLICANT: Glucksmann, Maria Alexandra
TITLE OF INVENTION: 31939, A NOVEL, HUMAN LEUCINE-RICH REPEAT
            CURRENT APPLICATION NUMBER: PCT/US01/10462
CURRENT FILING DATE: 2001-03-30
PRIOR APPLICATION NUMBER: 09/784,748
PRIOR FILING DATE: 2001-02-14
PRIOR APPLICATION NUMBER: 09/649,167
PRIOR FILING DATE: 2000-08-23
PRIOR FILING DATE: 2000-08-23
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                                                                                                                                    APPLICANT: Liu, Chenghua
APPLICANT: Tang, Y. Tom
APPLICANT: Drmanac, Radoje T.
ITTLE OF INVENTION: METHODS AND MATERIALS RELATING TO INSULIN-LIKE GROWTH FACTOR BINITITLE OF INVENTION: PROTEIN-LIKE POLYPEPTIDES AND POLYNUCLEOTIDES
FILE REFERENCE: 21272-051
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SOFTWARE: FastSEQ for Windows Version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: 31939, A NOVEL HUMAN LEUCINE-RIGHTEL OF INVENTION: FAMILY MEMBER AND USES THEREOF
                                                                                                                                                                                                                                                         PPLICANT:
                                                                                                                                                                                                                                                                                              PPLICANT: HYSEQ,
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TYPE: PRT
ORGANISM: Artificial Sequence
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APPLICATION NUMBER: 09/540,217
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Asundi, Vinod
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                                                                                                                                                                                                                                                                       Victoria
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80.0%;
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Pred. No. 0.81;
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; ORGANISM: Homo sapiens
PCT-US01-11988-133
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-266-091-4
                                                                                                                                                                                                                                                                                                                                         PCT-US01-11988-133
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Best Local Similarity
Thes 7; Conserv.
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                                                 SEQ ID NO 133
LENGTH: 359
                                                                                                                                                                                                                                                                                                    Sequence 133, Application PC/TUS0111988 GENERAL INFORMATION:
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Best Local
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                                                                               PRIOR APPLICATION NUMBER: 60/199, 384
PRIOR FILING DATE: 2000-04-25
NUMBER OF SEQ ID NOS: 2267.
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                PRIOR APPLICATION NUMBER: 60/256, 931 PRIOR FILING DATE: 2000-12-21
                                                                                                                                                                                CURRENT APPLICATION NUMBER: PCT/US01/11988
CURRENT FILING DATE: 2001-01-12
PRIOR APPLICATION UMBER: 60/229, 358
PRIOR FILING DATE: 2000-04-12
                                                                                                                                                                                                                                                    APPLICANT: Human Genome Sciences, Inc.
TITLE OF INVENTION: Albumin Fusion Proteins
FILE REFERENCE: PF546PCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 08/657,753
PRIOR FILING DATE: 1996-05-30
NUMBER OF SEQ ID NOS: 10
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CURRENT FILING DATE: 1999-03-10
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APPLICANT: Semple, Sean C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 14
SOFTWARE: PatentIn version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Bennett, C. Frank
TITLE OF INVENTION: ENHANCED EFFICACY OF LIPOSOMAL ANTISENSE THERAPY
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                                                                                                                                                                                                                                                                                                                                                                                                        284 EDSGIYVCE 292
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Hope, Michael
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Pred. No. 11;
1; Mismatches
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Pred. No.
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Search completed: June 4, 2001, 12:21:29 Job time: 591 sec	QY 1 EDSGLYQCEAAT 12 :	Query Match 64.1%; Score 41; DB 1; Length 359; Best Local Similarity 66.7%; Pred. No. 8.3; Matches 8; Conservative 1; Mismatches 3; Indels
4, 2001,	14 2	64.1%; 66.7%; tive
12:21:29	•	%; Score 41; DB 1; %; Pred. No. 8.3; 1; Mismatches
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Result
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Maximum Match 100%
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Perfect score:
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Maximum DB seq length: 200000000
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                                                                                                                                                                                                                                             Score
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/cgn2_6/ptodata/2/paa/US098_COMB.pep:*
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/ptodata/2/paa/US093_COMB.pep:
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ptodata/2/paa/US087_COMB.pep:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   otodata/2/paa/US088_COMB.pep:
                                               3 US-60-160-209-4252
3 US-60-160-203-4414
6 US-09-245-764-7
1 US-09-702-021-14
PCT-US00-05882-1128
US-08-332-562-94
US-08-452-973-94
US-07-861-895-25
                                                                                                                                                                                                                                                                                                             SUMMARIES
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Sequence 7, Appli
Sequence 1128, Appl
Sequence 1128, Appl
Sequence 94, Appl
Sequence 94, Appl
Sequence 25, Appl
Sequence 27859, A
Sequence 22468, A
                                                                                                                                                                                                                                           Description
                                                                                                                                                                         Sequence 4252, Ap
Sequence 4414, Ap
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	36	3 3	. 36	36	36	36	36	36	36	36	. 36	36	36.5	37	37	37	37	37	37	37	37	37	37	37	37	37	37	37	37	37.5	38	39	42	42	
	59.0	9.0	9	9	9.	9.	9	9	9	9	9	9	59.8	60.7	60.7	60.7	60.7	60.7	60.7	60.7	60.7	60.7	60.7	60.7	60.7	60.7	60.7	60.7	0	_	.٧	63.9	8	68.9	
	358	260	217	213	189	189	181	181	132	132	132	56	366	1970	1970	1908	1908	1907	1354	1354	1354	. 1354	1354	1354	1087	397	168	143	124	σ	490	368	407	9	
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	US-09-206-647-1	-09-497-191-861	-US01-01349-	-US96-15098A-	70-767-5099	-270-767-	-US01-01349-	-01326-13	-08-911-503A-6	-08-911-50	-US97-14436-6	S-09-733-089-158	S-09-107-532-429	-60-173-464-1798	S-60-167-217-2213	-60-191-681-2876	-60-191-637-	-60-173-464-2830	-60-191-681-607	-60-191-637-780	-60-173-464-631	-60-171-627-6	-60-167-217-7	-09-619-049-41	-09-417-507-42680	-09-489-039A-1	7-217-787	9-134-000-429	-09-733-089-16	-09-450-969-754	-09-248-796-1616	-09-248-796-164	S-09-347-801-23	US-60-167-217-7942	
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	Ą,	ce 861, Ap	664, 4	e 11,	0	ce 3578	e 83	137, A	e 634	ce 634	634,	e 158	equence 4292, A	equence 1798	equence	equence 28768,	e 36817,	e 2830	e 6078,	e 7800,	e 6310,	e 660,	e 7900	æ	e 4268	e 13098,	e 7877,	æ	æ	e 7543,	e 161	equence 16492,	e 23, App	Sequence 7942, Ap	

ALIGNMENTS

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В
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                                                                                                                                                           ; NAME/KEY: VARIANT;
LOCATION: (1)...(96)
; OTHER INFORMATION: Xaa = Any Amino Acid
US-60-160-209-4252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-60-160-209-4252
                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 4646
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4252
LENGTH: 96
TYPE: PRT
ORGANISM: HUMAN
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 4252, Applic GENERAL INFORMATION: APPLICANT: BONAZZI,
                                                                   Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 12; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: 1
TITLE OF INVENTION: 2
TITLE OF INVENTION: 1
FILE REFERENCE: CLOOC
                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/60/160,209 CURRENT FILING DATE: 1999-10-19
79 EDGNVLKRSPEL 90
                     1 EDGNVLKRSPEL 12
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                                                                     Score 61; DB 23;
Pred. No. 0.00038;
; Mismatches 0;
                                                                                                     Length 96;
                                                                     Indels
                                                                     0
                                                                     Gaps
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LENGTH: 102
; TYPE: PRT
; ORGANISM: HUMAN
US-60-160-203-4414
                                                                                      US-09-702-021-14
                                                                                                                                                                                                                                                                                                                ; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-245-764-7
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                                                  Sequence 14, Application US/09702021 GENERAL INFORMATION:
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Best Local
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Best Local Similarity
Matches, 12; Conserv
APPLICANT: C. FRASER TITLE OF INVENTION: FAIL MOLECULES AND USES THEREOF FILE REFERENCE: 7853-217
                                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 6374
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4414
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CURRENT FILING DATE: 1999-10-19
                                                                                                                                                                                                                                                                                                                                                                                                   OFTWARE: PatentIn
                                                                                                                                                                                                                                                                                                                                                                                                                                  SARLIER APPLICATION NUMBER: 60/073,972
SARLIER FILING DATE: 1998-02-06
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: ISOLATED HUMAN SECRETED PROTEINS,
TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING HUMAN SECRETED PROTEINS AND
TITLE OF INVENTION: USES THEREOF
FILE REFERENCE: CL000116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PPLICANT: Maxwell, Kelly F.
PPLICANT: Garrett, Thomas P.J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ENERAL INFORMATION:
APPLICANT: BONAZZI, VIVIEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ILE REFERENCE: 4102-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PPLICANT: McKenzie,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PPLICANT: Hogarth, P. Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ITLE OF INVENTION: THREE DIMENSIONAL STRUCTURES AND MODELS OF FC RECEPTORS TILE OF INVENTION: AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                   ENGTH: 261
                                                                                                                                                           246
                                                                                                                                                                                                                                                                                                                                                                                     ID NO 7
                                                                                                                                                                                                                            Local Similarity
nes 12; Conserv
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                                                                                                                                                                       1 EDGNVLKRSPEL 12
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                                                                                                                                                         EDGNVLKRSPEL . 257
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/09/245,764 FILING DATE: 1999-02-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      [LING DATE: 1998-09-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Powell, Maree S. McKenzie, Ian F.C.
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                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CATION NUMBER:
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Best Local Similarity
""+" hes 12; Conserv
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                                                                                                                                                                                                                                                                                                             Matches
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SEQ ID NO 1128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ fo:
SEQ ID NO 14
LENGTH: 374
                                                                                                                                                                                            -08-332-562-94
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT FILING DATE: 2000-0: EARLIER APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EARLIER FILING DATE: 1999-03-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: PCT/US00/05882
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: Human Cancer Associated FILE REFERENCE: PA106PCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Craig Rosen, APPLICANT: Steve Ruben
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/702,021 CURRENT FILING DATE: 2000-10-30
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TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: SITE
                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE
                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: SITE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ENGTH:
                             CORRESPONDENCE ADDRESS
                                                                                                                                                 APPLICANT:
                                            NUMBER OF SEQUENCES:
                                                                                                     APPLICANT:
                                                                                                                                                                                                                                                    290 EDGNVLKRSPEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              265 EDGNVLKRSPEL 276
                                                                                                                                                                                                                                                                                                                            Local
                                                                                                                                                                                                                                                                                1 EDGNVLKRSPEL 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 EDGNVLKRSPEL 12
                                                                                                                                                                                                                                                                                                             l Similarity
12; Conser
                                                           INVENTION:
                                                                                                                                                                            Application US/08332562
: Foley & Lardner 3000 K Street, N.W.,
                                                                                    BONADONNA, Lisa
                                                       HULETT, Mark D.

ENTION: FC RECEPTOR FOR IMMUNOGLOBULIN
                                                                                                   SCOTT, Bernadette M
                                                                                                                  HIBBS, Margaret L.
                                                                                                                                              MCKENZIE,
                                                                                                                                  HOGARTH, Mark
                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Application
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                                                                                                                                                                                                                                                    301
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                                                                                                                                                                                                                                                                                                                                                                                                Xaa equals any of the
                                                                                                                                                                                                                                                                                                                         100.0%;
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                                                                                                                                            Ian F.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Windows Version
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PC/TUS0005882
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                                                                                                                                                                                                                                                                                                                           Score 61;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                naturally
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gene Sequences and Polypeptides
                                                                                                                                                                                                                                                                                                                                     Length 399;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 374;
                                                                                                                                                                                                                                                                                                                                                                                              occurring L-amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       L-amino acids
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STREET:

Suite 500

COUNTRY:

USA

20007-5109

CITY: Washington STATE: D.C.

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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 91 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                       quence 94, Applicat
ENERAL INFORMATION:
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MEDIUM TYPE: Floppy disk
                                                                                                  SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 54
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
      CLASSIFICATION DATA:
PRIOR APPLICATION NUMBER: US 08/332,562
                                                                                                                                                                                                                                                                                                     ORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                          APPLICANT
                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
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APPLICATION NUMBER: US 07/896,457
FILING DATE: 27-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                     UMBER OF SEQUENCES:
                                                  FILING DATE: 05
CLASSIFICATION:
                                                                                                                                  COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                      CITY: Washington STATE: D.C.
                                                                                                                                                                                                                  COUNTRY:
                                                                                                                                                                                                                                                                  STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          77 EDSSVLKRSPEL 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX:
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FILING DATE: 31-OCT-1994
                                                                                     APPLICATION NUMBER: US/08/462,973
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE:
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                                                                                                                                                                                                     20007-5109
                                                                                                                                                                                                                                                                                                                                         INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Application US/08462973
                                                                                                                                                                                                                                                                    3000 K Street, N.W., Suite 500
                                                                                                                                                                                                                     USA
                                                                                                                                                                                                                                                                                                                                                                                                    MCKENZIE, Ian F.C.
HOGARTH, Mark P.
HIBBS, Margaret L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (202)672-5399
                                                                                                                                                                                                                                                                                                                                    HULETT, Mark D.

TENTION: FC RECEPTOR FOR IMMUNOGLOBULIN
                                                                                                                                                                                                                                                                                                                                                                       BONADONNA, Lisa
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                                                                                                                                                                                                                                                                                     Foley & Lardner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Stephen A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               UMBER: US/08/332,562
31-OCT-1994
                                                                05-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                        Bernadette M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          82.0%;
83.3%;
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Pred. No.
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                                                                                                                    Version #1.30
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Best Local Similarity
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REFERENCE/DOCKET NUMBER: 16.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
TELEPAX: (202)672-5399
TELEPAX: (202)672-5399
                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 91 amino acids
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 'AU PJ 7045
FILING DATE: 25-OCT-1989
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 16-SEP-
PRIOR APPLICATION DATA:
                                                                                                   PRIOR APPLICATION DATA:
                                                                                                                                                                     CURRENT APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: HYBRID FC RECEPTOR MOLECULES NUMBER OF SEQUENCES: 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
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                                                                    APPLICATION NUMBER: PCT/AU90/00513 FILING DATE: 25-OCT-1990
                                                                                                                    FILING DATE: 19
CLASSIFICATION:
                                                                                                                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         77 EDSSVLKRSPEL 88
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                                                                                                                                                     APPLICATION NUMBER:
                                                                                                                                                                                    SOFTWARE:
                                                                                                                                                                                                                                                                                    COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: BENT, Stephen A REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
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                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Foley & Lardner
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                                                                                                                                                                                                                                                                                        USA
                                                                                                                                                                                                                                                                                                                                                                                                                     MCKENZIE, Ian F. C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                 PatentIn Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                        OSMAN, Narin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HOGARTH, Phillip M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          UMBER: AU PH6166
26-MAY-1986
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27-MAY-1992
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                                                                                                                                                     US/07/861,895
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Pred. No. 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                      Version #1.25
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Gaps

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Query Match
Best Local Similarity
Thes 6; Conserv
; ORGANISM: DROSOPHILA
US-60-191-681-22468
                                                                                                                                                                                                                                          RESULT 10
US-60-191-681-22468
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                                         SOFTWARE: Fast
SEQ ID NO 22468
LENGTH: 99
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JS-60-191-637-27859
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 27859
LENGTH: 99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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Best Local Similarity
                                                                                                     CURRENT APPLICATION NUMBER: US/60/191,681 CURRENT FILING DATE: 2000-03-23
                                                                                      NUMBER OF SEQ ID NOS:
                                                                                                                                          FILE REFERENCE:
                                                                                                                                                                                                               equence 22468, Application US/60191681 ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/60/191,637
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: GENE SEQUENCES EXPRESSED FROM FITLE OF INVENTION: GENE SEQUENCES EXPRESSED FROM FITLE OF INVENTION: HERE FITLE OF INVENTION: HERE FITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYLE OF INVENTION: GENE SEQUENCES EXPRESSED FROM THE DROSOPHILA GENOME, AND ITLE OF INVENTION: USES THEREOF ILE REFERENCE: CL000392
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS
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FELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 280 EDSSVLKRSPEL 291
                                                                                                                                                                                                                                                                                                      36 DDGNVIKKGPD 46
                                                                                                                                                                                                                                                                                                                      1 EDGNVLKRSPE 11 : |: |:
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                                                                          FastSEQ for
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                                                                                                                                                    USES THEREOF.
                                                                                                                                                                  ACID MOLECULES ENCODING
                                                                                                                                                                                ISOLATED DROSOPHILA PROTEINS,
                                                                                                                                                                                                                                                                                                                                                                              68.9%;
54.5%;
                                                                       Windows Version 4.0
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83.3%;
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                                                                                                                                                                                                                                                                                                                                                                              Score 42;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                Mismatches
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0.36;
                                                                                                                                                                                                                                                                                                                                                                                            DB 23;
                                                                                                                                                             ISOLATED DROSOPHILA PROTEINS AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1; Indels
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Query Match

68.9%;

Score 42;

DВ

23;

Length 99;

US-09-248-796-16492 : Sequence 16492, Application US/09248796 : GENERAL INFORMATION:

APPLICANT: Keith Weinstock et al TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBI

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RESULT 13

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ORGANISM: Arabidopsis thaliana US-09-347-801-23
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US-60-167-217-7942
                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-347-801-23
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                                                              Matches
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 7942
LENGTH: 299
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: L1, Peter W. D.
TITLE OF INVENTION: ISOLATED DROSOPHILA PROT
TITLE OF INVENTION: ACID MOLEUCLES ENCODING
TITLE OF INVENTION: THÉREOF
FILE REFERENCE: CL000152
                                                                                           Query Match
                                                                                                                                                                                                            SEQ ID NO 23
                                                                                                                                                                                                                                                                                                                                                                                    Sequence 23, Application US/09347801 GENERAL INFORMATION:
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Best Local
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Matches 6; Conserv
                                                                                                                                                                                                                                         EARLIER APPLICATION NUMBER: 60/092,438 EARLIER FILING DATE: July 10, 1998 NUMBER OF SEQ ID NOS: 23
                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/347,801 CURRENT FILING DATE: 1999-07-02
                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Allen, Steve APPLICANT: Lee, Jian Mi
                                                                                                                                                                                                                           SOFTWARE: Microsoft Office 97
                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Plant Protein Kinases FILE REFERENCE: BB-1171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/60/167,217
CURRENT FILING DATE: 1999-11-24
                                                                                                                                                                           LENGTH: 407
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       236 DDGNVIKKGPD 246
21
                                                                            Local
                            2 DGNVLKRSPEL 12
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DGDALKRRPEL 31
                                                             Similarity 72.
8; Conservative
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                                                                            68.9%;
72.7%;
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54.58;
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                                                           Score 42; DB Pred. No. 13; 1; Mismatches
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Pred. No.
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Mismatches
                                                                          DB 17;
13;
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                                                                                        Length 407;
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NUMBER OF SEQ ID NOS: 28206; SEQ ID NO 16167; LENGTH: 490; TYPE: PRT; ORGANISM: Candida albicans US-09-248-796-16167
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US-09-248-796-16167
; Sequence 16167, Application US/09248796
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Best Local Similarity
Thes 7; Conserve
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US-09-450-969-7543
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                                                                                                                                                                                          NUMBER OF SEQ I
SEQ ID NO 7543
LENGTH: 166
TYPE: PRT
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Best Local Similarity 50.0%;
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                  Sequence 7543, Application US/09450969
GENERAL INFORMATION:
APPLICANT: Lynn Doucette-Stamm et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCCUS
TITLE OF INVENTION: EPIDERMIDIS FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: PATH99-09A
                                                                     Matches
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Best Local Similarity
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                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/450,969
CURRENT FILING DATE: 1999-11-29
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CURRENT FILING DATE: 1999-02-12
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EQ ID NO 16492
LENGTH: 368
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CURRENT FILING DATE: 1999-02-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TLE REFERENCE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICAN ITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
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::||: | ||:
330 KNGNIFKLSPEI 341
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1:||::|| ::|| 109 EEGNIIKRKTGVLMLNPEL 127
                                  1 EDGNVLKR----,---SPEL 12
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58.3%;
                                                                 Score 37.5; D
Pred. No. 32;
4; Mismatches
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Pred. No.
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Pred. No.
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Search completed: June 4, 2001, 12:20:04 Job time: 517 sec

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Result
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Match
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Copyright (c) 1993 - 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /cgn2_6/ptodata/1/paa/US06_NEW_COMB.pep:*
/cgn2_6/ptodata/1/paa/US07_NEW_COMB.pep:*
/cgn2_6/ptodata/1/paa/US08_NEW_COMB.pep:*
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US-09-549-066-36
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US-09-739-449-11789

PCT-US01-04098A-1252

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PCT-US01-04098A-1283
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PCT-US01-04098A-3229
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US-09-739-449-11283
US-09-270-849B-18364
                     US-09-549-066-35
US-09-327-563B-4
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Compugen Ltd.
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US-09-284-107-27
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Best Local Similarity
Matches 12; Conserv
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ'ID NO 27
LENGTH: 12
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NUMBER OF SEQ ID NOS: 13351
SEQ ID NO 11283
                                       APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
TITLE OF INVENTION: Agrobacterium tumefaciens
FILE REFERENCE: 38-10(15490)C
CURRENT APPLICATION NUMBER: US/09/739,449
CURRENT FILING DATE: 2000-12-19
PRIOR APPLICATION NUMBER: US 09/514,000
PRIOR EILING DATE: 2000-02-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/284,107
CURRENT FILING DATE: 1999-10-25
PRIOR APPLICATION NUMBER: PCT/NL97/00557
PRIOR FILING DATE: 1997-10-07
PRIOR APPLICATION NUMBER: EP 96202791.8
PRIOR FILING DATE: 1996-10-08
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE: Oligopeptides representing the extracellular OTHER INFORMATION: Oligopeptides representing the extracellular OTHER INFORMATION: domain of CD64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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LENGTH: 124

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245 EDNQVWKQSPQL 256

1 EDGNVLKRSPEL 12

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RESULT 3
US-09-270-849B-183641
; Sequence 183641, Application US/09270849B
; GENERAL INFORMATION:
APPLICANT: Swimmer et al.
; TITLE OF INVENTION: Insect genome survey devices
; TITLE OF INVENTION: OF THE O
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         뮹
                                                                                                                                                                                                                                            ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCT-US01-11988-133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     : OTHER INFORMATION: Description of Artificial Sequence: Synthetic : OTHER INFORMATION: at any location having Xaa, Xaa means any amino acid US-09-270-849B-183641
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   밁
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Best Local Similarity
Yeatches 7; Conserve
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                                                                                                                                                                                                                                                                                                               SEQ ID NO 133
LENGTH: 359
TYPE: PRT
                                                                                                           Query Match
Best Local :
                                                                  Matches
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NUMBER OF SEQ ID NOS: 195450
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 183641
LENGTH: 189
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                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 60/199, 384
PRIOR FILING.DATE: 2000-04-25
NUMBER OF SEQ ID NOS: 2267
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: PCT/US01/11988 CURRENT FILING DATE: 2001-01-12 PRIOR APPLICATION NUMBER: 60/229, 358 PRIOR PILING DATE: 2000-04-12 PRIOR APPLICATION NUMBER: 60/256, 931 PRIOR APPLICATION NUMBER: 60/256, 931 PRIOR FILING DATE: 2000-12-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/270,849B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ITLE OF INVENTION: Albumin Fusion Proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PPLICANT: Human Genome Sciences, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Artificial Sequence FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LE REFERENCE: PF546PCT
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                                                                  Local Similarity
les 7; Conserv
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nes 7; Conserv
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                                                                  Conservative
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                                                                                                59.0%;
58.3%;
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                                                       Score 36; DB Pred. No. 73; 2; Mismatches
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Pred. No.
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34;
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US-09-270-849B-191599
; Sequence 191599, Application US/09270849B
; GENERAL INFORMATION:
; APPLICANT: Swimmer et al.
; TITLE OF INVENTION: Insect genome survey devices
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RESULT

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В
                                                                                                                             FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Synthetic US-09-270-849B-192159
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US-09-833-245-133
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                                                                                    Query Match
Best Local
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Best Local Similarity 58.1
Matches 7; Conservative
                                                                  Matches
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                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 192159, Application US/09270849B
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SEQ ID NO 133
LENGTH: 359
TYPE: PRT
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                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/270,849B
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 195450
                                                                                                                                                                                                                                                                                                                                              FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Swimmer et al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR FILING DATE: 2000-12-21
PRIOR APPLICATION NUMBER: 60/199, 384
PRIOR FILING DATE: 2000-04-25
NUMBER OF SEQ ID NOS: 2267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/833,245
CURRENT FILING DATE: 2001-04-12
                                                                                                                                                                                                  LENGTH: 193
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Human Genome Sciences, Inc. TITLE OF INVENTION: Albumin Fusion Proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR FILING DATE: 2000-04-12
PRIOR APPLICATION NUMBER: 60/256, 931
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               245 EDNQVWKQSPQL 256
                                                                                  Local
                1 EDGNVLKRSP 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 EDGNVLKRSPEL
EDGVVLRRPP 15
                                                           n 57.4%;
Similarity 70.0%;
7; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12
                                                                                                                                                                                                                                                                                                                                                         Insect genome survey devices
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                                                              Score 35; DB
Pred. No. 54;
1; Mismatches
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                                                                               DB 5;
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                                                                                               Length 193;
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                                                                Indels:
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Query Match
Best Local Similarity
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Conservative

1; Mismatches

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Indels

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Gaps

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57.4%; 70.0%;

Score 35; DB 5; Pred. No. 1.6e+02;

Length 470

; OTHER INFORMATION: ; OTHER INFORMATION: US-09-270-849B-191599

Description of Artificial Sequence: Synthetic at any location having Xaa, Xaa means any amino acid

ORGANISM: Artificial Sequence

SEQ ID NO 191599

LENGTH: 470

TYPE: PRT

FEATURE:

NUMBER OF SEQ ID NOS: 1 SOFTWARE: PatentIn Ver.

CURRENT FILING DATE: 1999-03-17

FILE REFERENCE:

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Query Match
Best Local Similarity
Thes 6; Conserv
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US-09-549-066-37
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                                                                                                         TOPOLOGY:
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                                                                                                                                                                                  NFORMATION FOR SEQ ID NO:
                                                                                                                                                  SEQUENCE CHARACTERISTICS
LENGTH: 497 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Re
                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 13-APRIL-2000 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                     FILING DATE: 24-JUN-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    IIILE OF INVENTION: Programmed Cell Death Genes and Proteins NUMBER OF SEQUENCES: 85
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          282 EDGVVLRRPP 291
                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 08/080,850 FILING DATE: 24-JUN-1993
                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: FILING DATE: 10-JU
                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/09/549,066
                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Washington
                                                                                                                                       TYPE:
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                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                             REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
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1 EDGNVLKRSPEL 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         USA
                               Conservative
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                                                                                                                       both
                                                                                                                                                                                                                                                                                                                                                                                                                                              PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                               (202) 371-2600
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                                              Score 35;
Pred. No.
                               Mismatches
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                                                         Length 497;
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Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                        RESULT 10
                                                   SOFTWARE: PatentIn Ver. SEQ ID NO 1066
                                                                                                                                                                                                       GENERAL INFORMATION:
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                                                                                                                                                                                                                       Sequence 1066, Application PC/TUS0101312
                                                                                                                   CURRENT APPLICATION NUMBER: PCT/US01/01312 CURRENT FILING DATE: 2001-05-09
                                                                                                                                                     APPLICANT: Human Genome Sciences, Inc., et al. TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies FILE REFERENCE: PTZ32PCT
                                                                                   Prior application data removed - refer to PALM or file wrapper NUMBER OF SEQ ID NOS: 1510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NFORMATION FOR SEQ
ORGANISM: Homo sapiens
                                   ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 10-UN-1994
APPLICATION NUMBER: US 08/080,850
FILING DATE: 24-JUN-1993
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION: TELEPHONE: (202) 371-2600
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                                                                                                                                                                                                                                                                                                            471 QGSNILKQMPEL 482
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465 QGANILKQMPEL 476
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FILING DATE: 13-APRIL-2000
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SOFTWARE: Patenti
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SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                              Score 35;
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                                                                                                                                                                                                                                                                                                                                                                                                                 DB 5;
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FEATURE:

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; ORGANISM: Agrobacterium tumefaciens US-09-739-449-12341
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; LOCATION: (32)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
PCT-US01-01312-1066
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Best Local Similarity bu...
Conservative
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SEQ ID NO 12341
LENGTH: 216
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Best Local Similarity
Matches 7; Conser
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APPLICANT: Slater, Steven C.
TITLE OF INVENTION: Agrobacterium tumefaciens Genome Sequences
FILE REFERENCE: 38-10(15490)C
CURRENT APPLICATION NUMBER: US/09/739,449
CURRENT FILING DATE: 2000-12-19
PRIOR APPLICATION NUMBER: US 09/514,000
PRIOR APPLICATION UBBE: US 09/514,000
PRIOR APPLICATION UBBE: US 09/514,000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 12341, Application US/09739449
              PRIOR APPLICATION NUMBER: 09/560,875
PRIOR FILING DATE: 2000-04-27
PRIOR APPLICATION NUMBER: 09/496,914
PRIOR FILING DATE: 2000-02-03
                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: PCT/US01/04098A CURRENT FILING DATE: 2001-02-05 PRIOR APPLICATION NUMBER: NOT YET Assigned PRIOR FILING DATE: 2001-01-30 PRIOR APPLICATION NUMBER: 09/728,422 PRIOR APPLICATION NUMBER: 09/693,325 PRIOR APPLICATION NUMBER: 09/693,325
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 equence 1261, Application PC/TUS0104098A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      -US01-04098A-1261
                                                                                             RIOR APPLICATION NUMBER: 09/598,075
RIOR FILING DATE: 2000-06-20
                                                                                                                                     RIOR FILING DATE: 2000-07-19
                                                                                                                                                                          RIOR APPLICATION NUMBER: 09/654,936
RIOR FILING DATE: 2000-09-01
                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 09/663,561
PRIOR FILING DATE: 2000-09-15
                                                                                                                                                                                                                                                            RIOR FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                ITLE OF INVENTION: Novel nucleic Acids and Polypeptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     180 GNWLKRSP 187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3 GNVLKRSP 10
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87.5%;
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Pred. No.
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Pred. No.
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35;
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PCT-US01-11797-29

Sequence 29, Application PC/TUS0111797 GENERAL INFORMATION:

APPLICANT: SMITHKLINE BEECHAM CORPORATION
APPLICANT: SMITHKLINE BEECHAM p.l.c.
TITLE OF INVENTION: NOVEL COMPOUNDS
FILE REFERENCE: GP50022
CURRENT APPLICATION NUMBER: PCT/US01/11797
CURRENT FILING DATE: 2001-04-11

PRIOR APPLICATION NUMBER: 60/196,603

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                                                                                                                                                     ; ORGANISM: Homo sapiens PCT-US01-04098A-3229
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PCT-US01-04098A-3229
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Best Local Similarity 4,...
Conservative
                                                                       Matches
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SEQ ID NO 3229
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                                                                                                    Query Match
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SEQ ID NO 1261
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: PCT/US01/04098A CURRENT FILING DATE: 2001-02:05
                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 3960
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: Not Yet Assigned PRIOR FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: Novel Nucleic Acids and Polypeptides FILE REFERENCE: 21272-029
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 09/728,422
                                                                                                                                                                                                            LENGTH: 232
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118 EDGSVLKQFLSETEKMSPE 136
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 109 EDGSVLKQFLSETEKMSPE 127
                                                                      Local Similarity hes 9; Conserva
                                   1 EDGNVLKR-----SPE 11
                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 2000-09-01
APPLICATION NUMBER: 09/620,325
FILING DATE: 2000-07-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 09/693,325
FILING DATE: 2000-10-20
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                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 09/560,875
FILING DATE: 2000-04-27
                                                                                                                                                                                                                                                                                                                                              FILING DATE: 2000-06-20
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                                                                      Conservative
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47.4%;
                                                                                  Score 34; DB 1;
Pred. No. 1e+02;
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Pred. No.
                                                                      Mismatches
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                                                                                                    Length 232;
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RESULT 15
US-09-739-449-11837
Sequence 11837, Application US/09739449
GENERAL INFORMATION:
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
TITLE OF INVENTION: Agrobacterium tumefaciens Genome Sequences and Uses Thereof
FILE REFERENCE: 38-10(15490)C
CURRENT APPLICATION NUMBER: US/09/739,449
CURRENT FILING DATE: 2000-12-19
PRIOR APPLICATION NUMBER: US 09/514,000
PRIOR FILING DATE: 2000-02-23
NUMBER OF SEQ ID NOS: 13351
SEQ ID NO 11837
LENGTH: 184
TYPE: PRI
ORGANISM: Agrobacterium tumefaciens
US-09-739-449-11837
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR FILING DATE: 2000-04-13
: PRIOR APPLICATION NUMBER: 60/199,417
: PRIOR FILING DATE: 2000-04-24
: NUMBER OF SEQ ID NOS: 48
: SOFTWARE: FASTSEQ for Windows Version 3.0
: SEQ ID NO 29
: LENGTH: 801
: TYPE: PRT
: ORGANISM: Homo sapiens
PCT-US01-11797-29
Search completed: June 4, 2001, 12:21:29
Job time: 591 sec
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Best Local Similarity 66.3
Matches 6; Conservative
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Best Local Similarity 54.5%;
Matches 6; Conservative
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599 DDGHVMSCSPE 609
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133 DGLILKRGP 141
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Pred. No. 4.7e+02;
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Pred. No. 1.
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Post-processing: Minimum Match 0%
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Listing first 45 summaries
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Maximum DB seq length: 2000000000
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                                                                                                                                                                                                                                                           Database
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Gapop 10.0 , Gapext 0.5
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60
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

No.	Score	Match	Score Match Length DB ID	B. B.	ID	Description
ш	60	100.0	374	21	US-09-702-021-14	Sequence 14, App
2	60	100.0	399	_	PCT-US00-05882-1128	Sequence 1128, Ap
ω	43	71.7	165	_	PCT-US01-03080-9	Sequence 9, Ap
4	43	71.7	195	ᆫ	PCT-US01-03080-8	Sequence 8, Appli
ر.	39	65.0	214	23	US-60-258-251-43	Sequence 43, App
6	39	65.0	338	18	US-09-467-008-6	Sequence 6, Appl
7	39	65.0	380	ω	US-07-922-466-6	Sequence 6, Appli
8	39	65.0	380	ω	US-07-922-466A-6	Sequence 6, Appli
9	39	65:0	380	7	US-08-394-442-6	Sequence 6, Ap
10	39	22	380	20	US-08-416-478-6	Sequence 6, Appli

· Ę	. 45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29 .	28	27 .	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12	1	
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٠.	175	165	158	157	128	128	127	73	73	73	73	264	610	581	577	515	514	145	131	126	498	498	498	498	498	498	471	471	471	470	470	470	422	380	380	
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•	60-177	-60-177-670-18	-09-107-532-46	-60-170-374-32	-60-182-485-	2-485-	-733-089-	924-695-	-706-7418	-08-706-741-5	T-US97-15872-5	-09-201	-09-273	-09-273-420-	-09-684-458-	09-273	-09-273-420-	60-197-873-177	60-147-499-	S-60-197-873-146	-08-474-988A	-08-474-988-	-08-416-478	-08-394-442	-07-92	-07-922	-08-4	-08-474	-08-394	-08-474-	US-08-474-988-7	-394-	US-09-467-008-4	-08-474-9	US-08-474-988-6	
	133,	e 182, A	4621,	3209	58,	57, Ap	3657	50,	50,	e 50,	50, Appl	684	4,	10,	e 2,	e 6,	Œ	e 177	4839,	ce 14	e 9,	e 9,	e 2,	e 9,	e 2,	e 2,	e 2,	e 2	2	7,	7,	7,	e 4,	6	•	

ALIGNMENTS

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                             PCT-US00-05882-1128
; Sequence 1128, Application PC/TUS0005882
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                 ; ORGANISM: Homo sapiens US-09-702-021-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 1
US-09-702-021-14
                                                                             RESULT
                                                                                                                                                                                                                                                                                                                             SOFTWARE:
SEQ ID NO 14
                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 12; . Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 14, Application US/09702021
                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Craig Rosen, APPLICANT: Steve Ruben
                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/702,021
CURRENT FILING DATE: 2000-10-30
                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: C. Fraser
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                           FILE REFERENCE:
                                                                                                                                                                                                                                                                                                LENGTH: 374
TYPE: PRT
                                                                                                                         277 ELQVLGLQLPTP 288
                                                                                                                                         1 ELQVLGLQLPTP 12
                                                                                                                                                                                                                                                                                                                                              FastSEQ for Windows Version 3.0
                                                                                                                                                                                       . Conservative
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                                                                                                                                                                                                     Score 60; DB 21; Length 374; Pred. No. 0.073;
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                                                                                                                                                                                       Indels
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RESULT 4
PCT-US01-03080-8
; Sequence 8, App
                                                                             Db .
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                                                                                                                                                                                                                             PCT-US01-03080-9
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LENGTH: 399
                                                                                                                                                             Query Match
Best Local
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Best Local Similarity 100.0%;
Matches 12; Conservative
                                                                                                                                              Matches
                                                                                                                                                                                                                                       NAME/KEY: SITE LOCATION: (126)
OTHER INFORMATION: X
NAME/KEY: SITE LOCATION: (158)
OTHER INFORMATION: X
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EARLIER APPLICATION NUMBER: 60/124,270 EARLIER FILING DATE: 1999-03-12 NUMBER OF SEQ ID NOS: 1694
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: PCT/US00/05882
                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: \15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: PCT/US01/03080 CURRENT FILING DATE: 2001-01-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: Human Cancer Associated Gene Sequences and Polypeptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TILE REFERENCE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
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les 8; Conserv
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                                                                                                             1 ELQVIGLQLPTP 12
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                                                                             ELEVLGRELPVP 47
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Application PC/TUS0103080
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"ENTION: Bcl-2-like Polynucleotides, Polypeptides, and Antibodies
                                                                                                                                            Conservative
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                                                                                                                                                           71.78;
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Pred. No.
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RESULT
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                                                                    US-09-467-008-6; Sequence 6, Application US/09467008; GENERAL INFORMATION:
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Best Local S
Matches 8
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LENGTH: 214
TYPE: PRT
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Matches 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/60/258,251
CURRENT FILING DATE: 2000-12-27
NUMBER OF SEQ ID NOS: 102
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: ISOLA TITLE OF INVENTION: NUCL. TITLE OF INVENTION: AND FILE REFERENCE: CL0001039
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PRIOR APPLICATION NUMBER: 60/1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: PCT/US01/03080 CURRENT FILING DATE: 2001-01-31
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: PRIOR FILING DATE: 2000-0
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                                                                                                                                                                                                                                                                                                               LOCATION: (1)...(214)
OTHER INFORMATION: Xaa = Any Amino Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                               FEATURE: NAME/KEY: VARIANT LOCATION: (1)...(
                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: HUMAN
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                                                                                                                                                                  48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 ELQVLGLQLPTP
          ROMAGNANI, Serg
                                                           APPLICANT: TRIEBEL, Frederic
                                                                                                                                                                  ELRVLGKDLPRP 59
                                                                                                                                                                                              ELQVLGLQLPTP 12
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8; Conserv
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OF SEQUENCES:
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                                          MASTRANGEL1,
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66.7%;
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66.7%;
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                             Sergio
                                           Renato
                                                                                                                                                                                                                                           Score 39;
Pred. Nó.
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Pred. No.
              SPLICE VARIANTS
                                                                                                                                                                                                                             Mismatches
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24;
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CORRESPONDENCE ADDRESS

ADDRESSEE: BROWDY AND NEIMARK, P.L.L.C. STREET: 624 Ninth Street, N.W., Suite

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk

COUNTRY: USA

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UKGANISM: Homo sapiens;
SEQUENCE DESCRIPTION: SEQ ID NO: 6:
US-09-467;008-6
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Best Local Similarity
Matches 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                           quence 6, Application US/07922466 ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: 6: SEQUENCE CHARACTERISTICS:
                                       SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/922,466
                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PM PC COMPATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
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                                                                                                                                                                                                                                                                                                                UMBER OF SEQUENCES:
                                                                                                                                                                                                                                                       STREET:
                                                                                                                                                                                         COUNTRY:
                                                                                                                                                                                                                                                                      ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2 LQVLGLQLPTP 12
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MOLECULE TYPE: protein
ORIGINAL SOURCE:
ORGANISM: Homo sap
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/09/467,008 FILING DATE: 20-Dec-1999 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION: TELEPHONE: 202-628-5197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION: NAME: YUN, Allen C.
                                                                                                                                                                             20036
                                                                                                                                                                                                                 Washington : D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: EP 97401404.5
FILING DATE: 18-JUN-1997
APPLICATION NUMBER: PCT/EP98/03307
FILING DATE: 03-JUN-1998
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                                                                                                                                                                                                                                                     E: Sterne, Kessler, Goldstein & Fox 1225 Connecticut Avenue
ATION:
                                                                                                                                                                                             U.S.A.
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                         19920710
                                                                                                                                                                                                                                                                                                                            Lymphocytes, DNA Sequences Encoding These Proteins And Their Pharmaceutical And Biological Uses
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Pred. No. 1.9e+02;
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257 LTVLGLEPPTP 267

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                                                                          US-07-922-466A-6
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US-07-922-466A-6
 Matches
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             Query Match
Best Local Similarity
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GENERAL INFORMATION:
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Best Local Similarity
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INFORMATION FOR SEQ ID NO: 6:
                                                                                                                                              INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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LENGTH: 380 amino acids
TYPE: AMINO ACID
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                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 07
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)371-2600
                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                   MOLECULE TYPE:
                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION: NAME: Goldstein, Jorge A
                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
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                                                                                                                                                                                                                                                                             APPLICATION NUMBER: FILING DATE: 10-JUL CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: 1100 New CITY: Washington
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                                                                                                                                                                                                                                    NAME: Goldstein, Jorge A REGISTRATION NUMBER: 29,
                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
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                                                                                                                                ENGTH:
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                                                                                                                              380 amino acids
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Conservative
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                                                                                                                                                                         (202) 371-2540
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               65.0%;
72.7%;
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1; Mismatches
               Score 39; DB 3;
Pred. No. 2.2e+02;
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Pred. No. 2.2e+02;
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US-08-394-442-6
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GENERAL INFORMATION:
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Best Local Similarity
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NFORMATION FOR SEQ ID NO:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
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TITLE OF INVENTION:
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NAME: BROWDY, ROGER L.
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                                                                                                                                                                                                                                                                                                                        CITY: Washington
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Triebel,
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Pred: No. 2.2e+02;
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Best Local Similarity
"Arches 8; Conserva
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    08-474-988-6
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[NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
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ATTORNEY/AGENT INFORMATION:
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                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 08-SEP-
                                                                                                                                                                                                                                                                          SOFTWARE: Patentin Rel CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION:
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                                                                                     ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION:
REFERENCE/DOCKET NUMBER: HERCEND=1B TELECOMMUNICATION INFORMATION: TELEPHONE: (202) 628-5197
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER:
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SOFTWARE: PatentIn Release #1.0, Version #1.25
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                                                                                                                                                                      FILING DATE: 04-APR-1995
                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                   REGISTRATION NUMBER:
                                                                                                                                                                                     APPLICATION NUMBER:
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                                                                                                                                                        CLASSIFICATION:
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                                                                       YUN, Allen C.
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                                                                                                                                                                                                                                                                                                                                                                                                                                             419 Seventh Street, N.W., Suite 400
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                                                                                                                                                                                                                                                                                                                                          Floppy disk '
                                                                                                                                                                                                                                      UMBER: US/08/474,988
07-JUN-1995
                                                                                                      JMBER: US 07/854,644
08-SEP-1992
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72.7%;
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LYMPHOCYTES, DNA SEQUENCES ENCODING THESE PROTEINS AND
THEIR PHARMACEUTICAL AND BIOLOGICAL USES
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Pred. No. 2.2e+02;
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US-08-474-988A-6
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Best Local Similarity
Matches 8; Conserv
            Query Match
Best Local Similarity
 Matches
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                                                                                                                                                                             NFORMATION FOR SEQ ID NO:
                                                                                                   TYPE: amino acids
TYPOLOGY: line

(OLFORD
                                                                                                                                                                                                            REGISTRATION NUMBER: 37,971
REFERENCE/DOCKET NUMBER: HE
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 628-5197
                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/854,644

FILING DATE: 08-SEP-1992

ATTORNEY/AGENT INFORMATION:
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CURRENT APPLICATION DATA:
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                                                                                                  MOLECULE TYPE:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: BROWDY AND NEIMARK, P.L.L.C.
STREET: 419 Seventh Street, N.W., Suite 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US 08/416,478 FILING DATE: 04-APR-1995
                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM:
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amino acid
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                                                                                                                                                                                                                                                                                 YUN, Allen C.
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 Conservative
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SYSTEM: PC-DOS/MS-DOS
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                                                                                                  protein
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Pred: No. 2.2e+02;
                 Score 39;
Pred. No. 2
 Mismatches
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               DB 8;
2.2e+02;
                             Length 380;
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 Indels
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Gaps
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2 LQVLGLQLPTP 12

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ORGANISM: Homo sapiens ; SEQUENCE DESCRIPTION: SEQ ID NO: 4: US-09-467-008-4
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                                                                                                                                  US-08-394-442-7
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Best Local Similarity
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                                                                                              Sequence 7, Application:
                                                                                                                                                                                                                                                                                        Matches
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                    TITLE OF INVENTION:
                                                            APPLICANT:
                                                                            APPLICANT:
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                                                                                                                                                                                                                235 LTVLGLEPPTP 245
                                                                                                                                                                                                                                                2 LOVIGIQUETE 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: EP 97401404.5 FILING DATE: 18-JUN-1997 APPLICATION NUMBER: PCT/EP98/03307 FILING DATE: 03-JUN-1998 ATTORNEY/AGENT INFORMATION:
NAME: YUN, Allen C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/09/467,008 FILING DATE: 20-Dec-1999 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
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TITLE OF INVENTION: LAG-3 SPLICE VARIANTS
NUMBER OF SEQUENCES: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentln Release #1.0, Version #1.30
                                                                                                                  Application US/08394442
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REGISTRATION NUMBER: 37.971
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 422 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: 202-628-5197
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                                                          Triebel,
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                                                                            Hercend,
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                                                                            Thierry
                                                                                                                                                                                                                                                                                                          65.0%;
72.7%;
Lymphocytes, DNA Sequences Encoding These Proteins And Their Pharmaceutical And Biological Uses
                                    New Proteins Produced By Human
                                                            Frederic
                                                                                                                                                                                                                                                                                      Score 39; DB 10, pred. No. 2.4e+02;
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STREET: 415
STREET: Washington

ADDRESSEE:

E: BROWDY AND NEIMARK, P.L.L.C. 419 Seventh Street, N.W., Suite 400

COUNTRY:

U.S.A.

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Best Local Similarity 72.7%;
Matches 8; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:

NAME: BROWDY, ROGET L.

REGISTRATION UNMBER: 25,618
REFERENCE/DOCKET NUMBER: HERC
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 628-5197
TELEPHONE: (202) 737-3528
                    APPLICATION NUMBER: US 08
FILING DATE: 04-APR-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                            APPLICATION NUMBER: US/08/474 988 ETLING DATE: 07-UUN-1995 CLASSIFICATION 435 PRIOR APPLICATION DATA: APPLICATION NUMBER: US 08/416/478 APPLICATION NUMBER: US 08/416/478
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NFORMATION FOR SEQ ID NO:
                                                                                                                                                                    SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
COPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS: LENGTH: 470 amino acids
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/394,442
FILING DATE: 24-FEB-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION:
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APPLICATION NUMBER: US 07/854,644
                                                                                                                                                                                                                                                                                                 COUNTRY: UZIP: 20004
                                                                                                                                                                                                                                                                                                                                   CITY: Washington STATE: D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS
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419 Seventh Street, N.W., Suite 400
                                                                                                                                                                                                                                                                                                               U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hercend, Thierry
Triebel, Frederic
                                                                                                                                                                                                                                                                                                                                                                                                                                                LYMPHOCYTES, DNA SEQUENCES ENCODING THESE PROTEINS' AND THEIR PHARMACEUTICAL AND BIOLOGICAL USES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LYMPHOCYTES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NEW PROTEINS PRODUCED BY HUMAN
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Pred. No. 2
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Thes 8; Conserve
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                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 470 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                           NAME: YUN, Allen C.
REGISTRATION NUMBER: 37,971
REFERENCE/DOCKET NUMBER: HERCEND-1B
TELECOMMUNICATION INFORMATION:
229 LTVLGLEPPTP 239
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                         2 LQVLGLQLPTP 12
                                                     Conservative
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202) 737-3528
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Pred. No.
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Search completed: June 4, 2001, 12:20:05 Job time: 518 sec

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1: /cgn2_6/ptodatta/1/paa/PCT_NEW_COMB.pep:*
2: /cgn2_6/ptodatta/1/paa/US06_NEW_COMB.pep:*
3: /cgn2_6/ptodatta/1/paa/US07_NEW_COMB.pep:*
4: /cgn2_6/ptodatta/1/paa/US08_NEW_COMB.pep:*
5: /cgn2_6/ptodatta/1/paa/US08_NEW_COMB.pep:*
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Copyright (c) 1993 - 2000 Compugen Ltd.
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5 US-09-270-849B-180083

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PCT-US01-11988-2090
US-09-833-245-2090
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Sequence 4275, Ap	Sequence 184149,	Sequence 893, App	Sequence 893, App	Sequence 369, App	Sequence 1177, Ap	Sequence 17, Appl	Sequence 797, App	Sequence 181560,	Sequence 1225, Ap	Sequence 3676, Ap	Sequence 1708, Ap	Sequence 9005, Ap	Sequence 4, Appli	Sequence 12091, A	Sequence 882, App	Sequence 8808, Ap	Sequence 1275, Ap	

ALIGNMENTS

US-09-284-107-28

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Sequence 28, Application US/09284107
GENERAL INFORMATION:
APPLICANT: Logtenberg, Ton
APPLICANT: Logtenberg, Ton
APPLICANT: Logtenberg, Ton
APPLICANT: OF KULLEY
TITLE OF INVENTION: METHODS AND MEANS FOR SELE
TITLE OF INVENTION: AND PROTEINS HAVING SPECI
FILLE REFERENCE: 313632000600
CURRENT APPLICATION NUMBER: US/09/284,107
CURRENT FILING DATE: 1999-10-25
PRIOR APPLICATION NUMBER: PCT/NL97/00557
PRIOR APPLICATION NUMBER: EP 96202791.8
PRIOR APPLICATION NUMBER: EP 96202791.8
PRIOR FILING DATE: 1996-10-08
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Best Local S
Matches 12
SOFTWARE: PatentIn Ver. 2.0 SEQ ID NO 1035 LENGTH: 141
                                                                                                                                                                                                                              Sequence 1035, Application PC/TUS0101312 GENERAL INFORMATION:
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SOFTWARE: PASTSEQ for Windows Version 4.0
SEQ ID NO 28
LENGTH: 12
TYPE: PRT
                                                                                                                APPLICANT: Human Genome Sciences, Inc., et al. TITLE OF INVENTION: Nucleic Acids, Proteins, a FILE REFERENCE: PTZ32PCT CURRENT APPLICATION NUMBER: PCT/US01/01312 CURRENT FILING DATE: 2001-05-09
                                                                       Prior application data removed - refer to PALM or file wrapper NUMBER OF SEQ ID NOS: 1510\,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Artificial Sequence
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FEATURE: ORGANISM:

Homo sapiens

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; OTHER INFORMATION:
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; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids.
PCT-US01-01339-2813
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                                                                                                                                                                                   SEQ ID NO 1005
                                                                                                                                                                                                                                                                       Sequence 1005, Application PC/TUS0101329
GENERAL INFORMATION:
APPLICANT: Human Genome Sciences, Inc., et al
TITLE OF INVENTION: Nucleic Acids, Proteins,
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SOFTWARE: PatentIn Ver. 2.0
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                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: PCT/US01/01329
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Human Genome Sciences, Inc., et a TITLE OF INVENTION: Nucleic Acids, Proteins, FILE REFERENCE: PC006PCT
                                                                                                                                     ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Ver.
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                                                                                        NAME/KEY: SITE LOCATION: (1)
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OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
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Best Local Similarity

Query Match

56.7%;

Score 34;

DB 5;

Length 89;

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Best Local Similarity
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; LOCATION: (87)
; OTHER INFORMATION:
PCT-US01-11988-2091
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                                                                                                                                          SEQ ID NO 2091
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GENERAL INFORMATION:
OTHER INFORMATION: -09-833-245-2091
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                                                                                                                                                           SOFTWARE: PatentIn Ver.
                                                                                                                                                                           PRIOR FILING DATE: 2000-04-25
NUMBER OF SEQ ID NOS: 2267
                                                                                                                                                                                              PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Human Genome Sciences, Inc.
TITLE OF INVENTION: Albumin Fusion Proteins
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                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 60/199, 384
                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 60/256, 931
                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: PF546PC1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT FILING DATE: 2001-01-12
PRIOR APPLICATION NUMBER: 60/229, 358
PRIOR FILING DATE: 2000-04-12
                                                                                      ORGANISM:
                                                                                                                                                                                                                                                                RIOR FILING DATE:
                                                                                                                                                                                                                                                                               RIOR APPLICATION NUMBER: 60/229,
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                                                     NAME/KEY:
                                                                    FEATURE:
                                                                                                         TYPE: PRT
                                                                                                                                                                                                                                 RIOR FILING DATE:
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                                   LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Homo sapiens
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                                                                                                                        ENGTH:
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                                 SITE
(87)
                                                                                    Homo sapiens
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              Xaa equals any of the naturally occurring L-amino acids
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Pred. No.
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PCT-US01-01339-3249
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                                                                                          NAME/KEY: SITE
LOCATION: (116)
OTHER INFORMATION: X
'T-US01-11988-2090
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Best Local :
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APPLICANT: Human Genome Sciences,
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SOFTWARE: PatentIn Ver. 2.0
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ENERAL INFORMATION:
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CURRENT FILING DATE: 2001-01-12
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FITLE OF INVENTION: Albumin Fusion Proteins
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                                                                                                                                                     NAME/KEY: SITE
LOCATION: (109)
OTHER INFORMATION: Xaa
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                                                                                                                                                                                                                       NAME/KEY: SITE LOCATION: (107)
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                             Score 34; DB 1;
Pred. No. 1.1e+02;
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                                            Length 127;
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SEQ ID NO 2090
LENGTH: 127
                                                                                                                              NUMBER OF SEQ ID NOS: 13351
SEQ ID NO 9638
LENGTH: 335
TYPE: PRT
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                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 09/514,000 PRIOR FILING DATE: 2000-02-23
                                                                                                                                                                                                                            CURRENT FILING DATE: 2000-12-19
                                                                                                                                                                                                                                                            APPLICANT: Slater, Steven C.
TITLE OF INVENTION: Agrobacterium tumefaciens Genome Sequences
FILE REFERENCE: 38-10(15490)C
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CURRENT FILING DATE: 2001-04-12
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                                                                                                           ORGANISM: Agrobacterium tumefaciens
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LOCATION: (109)
OTHER INFORMATION:
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LOCATION: (116)
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)S: 2267
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Pred. No.
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Pred. No.
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                                                                                                         ; ORGANISM: Agrobacterium tumefaciens US-09-739-449-13150
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PCT-US01-04098A-1405
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SEQ ID NO 13150
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SEQ ID NO 1405
LENGTH: 350
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                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: US 09/514,000 PRIOR FILING DATE: 2000-02-23
                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/739,449
CURRENT FILING DATE: 2000-12-19
                                                                                                                                                                                                                                                                            APPLICANT: Slater, Steven C.
TITLE OF, INVENTION: Agrobacterium tumefaciens Genome Sequences and Uses Thereof
FILE REFERENCE: 38-10(15490)C.
                                                                                                                                                                                                                                                                                                                                                                               equence 13150
                                                                                                                                                                                                                                                                                                                               APPLICANT: Hinkle, Gregory J. APPLICANT: Slater, Steven C.
                                                                                                                                                TYPE: PRT
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                                                                                                                                                               ENGTH: 843
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APPLICATION NUMBER: Not Yet Assigned
FILING DATE: 2001-01-30
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APPLICATION NUMBER: 09/693,325
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 09/560,875
FILING DATE: 2000-04-27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LING DATE: 2000-09-01
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                  Conservative
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                                                                                                                                                                                                                                                                                                                                                                                 Application US/09739449
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2000-09-15
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                                 56.7%;
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                                 Score 34;
Pred. No. 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 34; DB 1; Le
Pred. No., 3.2e+02;
4; Mismatches 2;
                Mismatches
                                                  DB 5;
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                                                Length 843;
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; ORGANISM: Homo sapiens US-09-809-391-328
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Best Local Similarity
Matches 7; Conser
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SEQ ID NO 328
LENGTH: 23
TYPE: PRT
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NUMBER OF SEQ ID NOS: 116
                                  EARLIER APPLICATION NUMBER: 07/964,589
EARLIER FILING DATE: 1992-10-21
EARLIER APPLICATION NUMBER: PV-709-92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      09-178-115-51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: 186 Human Secreted proteins FILE REFERENCE: PZ002P2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Prior application data removed - NUMBER OF SEQ ID NOS: 761
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/809,39: CURRENT FILING DATE: 2001-03-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Ruben et al.
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                                                                                                FILING DATE: 1993-12-30
                                                                                                             FILING DATE: 1994-06-15
APPLICATION NUMBER: 08/177,093
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/09/178,115
FILING DATE: 1998-10-23
                                                                                                                                                                                                                                APPLICATION NUMBER: 08/485,863
                                                                                                                                                                                                                                                    APPLICATION NUMBER: 08/485,862 FILING DATE: 1995-06-07
                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 08/486,756 FILING DATE: 1995-06-07
                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 08/485,049 FILING DATE: 1995-06-07
                                                                                                                                                     APPLICATION NUMBER: 08/260,190
                                                                                                                                                                                             APPLICATION NUMBER:
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                                                                                                                                                                             LING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    LING DATE: 1997-01-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pastorekova, Silvia
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1995-06-07
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77.8%;
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Pred. No.
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; LENGTH: 257 ; TYPE: PRT ; ORGANISM: HUMAN US-09-178-115-51

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Sequence 1196, Application PC/TUS0101332

GENERAL INFORMATION:
APPLICANT: Human Genome Sciences, Inc., et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PJZ02PCT
CURRENT FILLING DATE: 2001-05-09
Prior application data removed - refer to PALM or file wrapper
NUMBER OF SEQ ID NOS: 1249
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 1196
SEQ ID NO 1196
TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: SITE
LOCATION: (184)
OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
PCT-US01-01332-1196
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PCT-US01-01332-1196
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Search completed: June 4, 2001, 12:21:30 Job time: 592 sec
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Best Local Similarity 66.7%;
Matches 6; Conservative
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Best Local Similarity 66.7
Matches 6; Conservative
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46 LELLGFQLP 54
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45 VKGINLPTP 53
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Pred. No. 4.5e+02;
1; Mismatches 2;
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GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.

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Result
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Maximum Match 100%
Listing first 45 summaries
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US-09-252-691C-7770
US-09-489-039A-7598
US-09-107-433-2896
                                                                                                                                       US-09-702-021-14
PCT-US00-05882-1128
US-07-861-895-25
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Sequence 11, Appl
Sequence 25, Appl
Sequence 27, Appl
Sequence 7770, Ap
Sequence 7770, Ap
Sequence 7598, Ap
Sequence 2896, Ap
Sequence 6248, Ap
Sequence 11281, A
Sequence 11281, A
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,	585,	732,	62	409	4	Sequence 4, Appli	Sequence 7106, Ap	8	8488	36,	24,	e 20,	Sequence 20, Appl	Sequence 52083, A	e 3686			19241,	Sequence 1831, Ap	1832	2, App	Sequence 1833, Ap		e 220	13791,	768, A	6717,	5647		27248	400, Ap	2875	5082	Sequence 35611, A	2793	

ALIGNMENTS

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; SOFTWARE: FastSEQ for Wi
; SEQ ID NO 14
; LENGTH: 374
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-702-021-14
                               RESULT 2
PCT-US00-05882-1128
PCT-0500-05882-1128
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Sequence 1128, Application PC/TUS0005882
GENERAL INFORMATION:
APPLICANT: Craig Rosen,
                                                                                                                                                                  Query Match
Best Local Similarity
Matches 12; Conserv
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Pred. No. 0.0083;
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                                                                                                                                                                                              Length 374;
                                                                                                                                                                     Indels
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Craig Rosen, Steve Ruben

TITLE OF INVENTION: Human Cancer Associated Gene

Sequences and Polypeptides

REFERENCE: PA106PC

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В
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MARLIER FILING DATE: 1999-03-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: PCT/US00/05882
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: SITE LOCATION: (349)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AME/KEY: SITE
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                                                                                                      REFERENCE/DOCKET NUMBER: 17
RE
                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 25-OCT-PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: Patentin Rel
                                                                                                                                                                                                                                                                                                        TTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORRESPONDENCE ADDRESS
                                                                                                                                                                                                                             NAME: BENT, Stephen A REGISTRATION NUMBER: :
                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: PCT/A
                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: AU PJ 7045 FILING DATE: 25-OCT-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OPERÁTING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 60/124,270
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FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3000 K Street, N.W.,
                                                                       (202)672-5399
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OSMAN, Narin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PatentIn Release #1,0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MCKENZIE, Ian F. C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HULETT, Mark D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HOGARTH, Phillip M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         IERINO, Francesco L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Foley & Lardner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19920624
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Suite 500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Version #1.25
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; ORGANISM: Enterobacter cloacae US-09-252-691C-7770
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Best Local Similarity
"hehes 6; Conserv
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; MOLECULE TYPE:
US-07-861-895-25
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                                                                                                                                                                            SEQ ID NO 7770
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LENGTH: 370
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APPLICANT: Keith G.
TITLE OF INVENTION:
TITLE OF INVENTION:
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Best Local Similarity
Matches 10; Conserv
                                               Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION:

TITLE OF INVENTION:

FILE REFERENCE: 107
                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 7770, Application US/09252691C
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                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/252,691C CURRENT FILING DATE: 1999-02-18
                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: US PRIOR FILING DATE: 1998-02-18 NUMBER OF SEQ ID NOS: 11326
                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT FILING DATE:
NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                   PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/252,691B
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                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                               ENGTH: 370
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2 WFHVLFYLA 10
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                                                                                                                                                                                                                                                                                                                                                 Keith G. Weinstock et al. TENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ENTEROBACTE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Keith G.
                                Conservative
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                                                                                                                                                                                                                                                                                                                    ON: CLOACAE FOR DIAGNOSTICS 107196.135
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NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ENTEROBACTE
CLOACAE FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                               .998-07-24
                                                                                                                                                                                                                                                        1999-02-18
IBER: US 60/094,145
                                                 61.8%;
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                                                                                                                                                                                                                                us 60/074,787
                                               Score 42;
Pred. No.
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                                                                  16;
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                                                                 Length 370;
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131 WYHGIFYLA 139

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Query Match
Best Local Similarity
Thes 5; Conserve
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US-09-107-433-2896
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JS-09-489-039A-7598
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             S-09-107-433-2899
Sequence 2896, Application US/09107433
GENERAL INFORMATION:
APPLICANT: Lynn A Doucette-Stamm and David Bush
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID
SEQUENCES RELATING TO STREPTOCOCCUS PNEUMONIAE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS:
SEQ ID NO 7598
LENGTH: 410
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CURRENT FILING DATE: 2000-01-27
PRIOR APPLICATION NUMBER: US 60/117,747
PRIOR FILING DATE: 1999-01-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 equence 7598, Appli
ENERAL INFORMATION:
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ITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA
ITLE OF INVENTION: PNEUMONIAE FOR DIAGNOSTICS AND THERAPEUTICS
                                                                 INFORMATION FOR SEQ ID NO: 2896:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: CD/ROM ISO9660
                                             SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                  APPLICATION NUMBER: 60/ 085131
FILING DATE: May 12, 1998
APPLICATION NUMBER: 60/051553
FILING DATE: July 2, 1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/09/107,433 FILING DATE: 30-Jun-1998 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS
                                                                                                          TELECOMMUNICATION INFORMATION: TELEPHONE: (781)893-5007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
TYPE: amino acid
                                                                                                                                                  REGISTRATION NUMBER: 40,489
REFERENCE/DOCKET NUMBER: GTC-011
                                                                                                                                                                                                                                                                                                                                                                                                                OPERATING SYSTEM: <Unknown>
SOFTWARE: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: <Unknown>
                                                                                                                                                                                             NAME: Ariniello, Pamela Deneke
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: Massachusetts
                      LENGTH: 104 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: GENOME THERAPEUTICS CORPORATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ITY: Waltham
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45.5%;
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Pred. No. 1
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                                                                                                                        ; ORGANISM: Enterobacter cloacae US-09-252-691-11281
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; ORGANISM: S.epidermidis
US-09-450-969-6248
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Best Local Similarity 60.
Matches 6; Conservative
                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/252,691B
CURRENT FILING DATE: 1999-02-18
NUMBER OF SEQ ID NOS: 11324
SEQ ID NO 11281
LENGTH: 418
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SEQ ID NO 6248
LENGTH: 244
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Lynn Doucette-Stamm et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCC
TITLE OF INVENTION: EPIDERMIDIS FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                TITLE OF INVENTION:
TITLE OF INVENTION:
FILE REFERENCE: 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT FILING DATE: 1999-11-29
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                                                                                                                                                                                                                                                                                                                      APPLICANT:
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LOCATION: (B) LOCATION 1...104
SEQUENCE DESCRIPTION: SEQ ID NO: 2896:
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                                                                                                                                                                                                                                                                                Keith G. Weinstock et al.
VENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ENTEROBACTE
VENTION: CLOACAE FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Application US/09450969
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Pred. No. 94;
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; Sequence 27932, Application US/09417507
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US-09-417-507-27932
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Best Local Similarity 4...
5; Conservative
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APPLICANT: Homburger et al.
TITLE OF INVENTION: Nucleic acids and proteins
FILE REFERENCE: File Reference: 7326-094
                                   SEQ ID NO 35611
LENGTH: 239
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SEQ ID NO 11281
                                                                                        CURRENT APPLICATION NUMBER: US/09/270,767
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517
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TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ASPERGILLUS
TITLE OF INVENTION: FUMICATUS FOR DIAGNOSTICS AND THERAPEUTICS
                                                                       SOFTWARE: PatentIn Ver. 2.0
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ORGANISM: Drosophila melanogaster
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CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/094,145
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                                                                                                                                                                                                                     Application US/09270767
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1998-02-18
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RESULT 15
US-09-741-669-400
; Sequence 400, Application US/09741669
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US-09-417-507-28756
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Best Local Similarity 75.0
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CURRENT FILING DATE: 1999-10-14
NUMBER OF SEO ID NOS: 44312
SEQ ID NO 28756
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SEQ ID NO 50828
LENGTH: 239
TYPE: PRT
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CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517
                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: KEITH G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster FILE REFERENCE: File Reference: 7326-094
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NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ASPERGILLUS FUMIGATUS FOR DIAGNOSTICS AND THERAPEUTICS
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GENERAL INFORMATION:
APPLICANT: Forsyth, R. Allyn
APPLICANT: Ohlsen, Kari L.
APPLICANT: Ohlsens identified as required for TITLE OF INVENTION: Genes identified as required for TITLE OF INVENTION. Genes identified as required for TITLE OF INVENTION. DIGNAL COLL.
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/cgn2_6/ptodata/1/paa/PCT_NEW_COMB.pep:*
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Copyright (c) 1993 - 2000 Compugen Ltd.
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/cgn2_6/ptodata/1/paa/US08_NEW_COMB.pep:*
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US-09-270-849B-187144
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US-09-811-284-165
US-09-811-284-168
                        PCT-USO1-04098A-1516
PCT-USO1-04098A-3484
PCT-USO1-04098A-1952
PCT-USO1-11988-662
PCT-USO1-11988-663
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US-60-248-505-811
US-60-248-505-1121
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US-09-270-849B-182343
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183288,
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168, App
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1265, Ap
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Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence
995I,	3573,	1605, 1	13267,			195, Ap	1004, Ap	•	1990, 7	1990, Ap	968, Ap	192136,	184164,	3818, A	364, App	9314, A	9//, Ap

ALIGNMENTS

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RESULT 2
US-09-739-449-9245
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US-09-284-107-29
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SOFTWARE: Fasi
SEQ ID NO 29
FUNCTH: 12
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CURRENT APPLICATION NUMBER: US/09/739,449
CURRENT FILING DATE: 2000-12-19
PRIOR APPLICATION NUMBER: US 09/514,000
PRIOR FILING DATE: 2000-02-23
NUMBER OF SEQ ID NOS: 13351
SEQ ID NO 9245
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CURRENT FILING DATE: 1999-10-25
PRIOR APPLICATION NUMBER: PCT/NL97/00557
                                                                                                                        APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
TITLE OF INVENTION: Agrobacterium tumefaciens
FILE REFERENCE: 38-10(15490)C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: EP 96202791.8 PRIOR FILING DATE: 1996-10-08
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TITLE OF INVENTION: METHODS AND MEANS FOR S
TITLE OF INVENTION: AND PROTEINS HAVING SP
FILE REFERENCE: 313632000600
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APPLICANT: de Kruif, Corne
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US-09-270-849B-183288
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                                                                                 Query Match
Best Local :
                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 13351
SEQ ID NO 11606
LENGTH: 331
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Best Local Similarity
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Best Local Similarity 63.6%;
Matches 7; Conservative
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LENGTH: 239
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                                                                                                                                                                                                                                                                               APPLICANT: Slater, Steven C.
TITLE OF INVENTION: Agrobacterium tumefaciens Genome Sequences
FILE REFERENCE: 38-10(1549)C
CURRENT APPLICATION NUMBER: US/09/739,449
CURRENT FILING DATE: 2000-12-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/270,849B
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 195450
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: US 09/514,000 PRIOR FILING DATE: 2000-02-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: Insect genome survey devices
                                                                                                                                                       TYPE: PRT ORGANISM: Agrobacterium tumefaciens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Swimmer et al.
                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Hinkle, Gregory J.
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202 VWFHMGFYTLI 212
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nilarity 54.5%;
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at any location having Xaa,
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Pred. No.
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; OTHER INFORMATION: US-09-270-849B-193851
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US-09-270-849B-185951
                                                                                                                                                                                                                        US-09-270-849B-182343
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NUMBER OF SEQ ID NOS: 195450
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 182343
LENGTH: 141
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CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 195450
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 185951
                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                      Sequence 182343, Application US/09270849B
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SEQ ID NO 193851
LENGTH: 61
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                                                                                    CURRENT APPLICATION NUMBER: US/09/270,849B CURRENT FILING DATE: 1999-03-17
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CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 195450
                                                                                                                                            TITLE OF INVENTION: Insect genome survey devices
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TITLE OF INVENTION: Insect genome survey devices
                                                                                                                          FILE REFERENCE:
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50.0%;
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Pred. No.
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Pred. No. 93;
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; OTHER INFORMATION:
; OTHER INFORMATION:
US-09-270-849B-182343

Description of Artificial Sequence: Synthetic at any location having Xaa, Xaa means any amino acid.

TYPE: PRT
ORGANISM: Artificial Sequence

FEATURE:

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1266 WFHFVFFI 1273

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; TYPE: PRT
; ORGANISM: Human
US-60-248-505-811
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Best Local Similarity
Fighter 5; Conserv
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US-60-248-505-811
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Best Local Similarity
                                           Query Match
Best Local Similarity
                                                                                                                                                                 SEQ ID NO 811
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CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 195450
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/60/248,505
CURRENT FILING DATE: 2000-11-15
                                                                                                                                                                                                                                        TITLE OF INVENTION: PROFILE REFERENCE: c1000918
                                                                                                                                                                               SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                              NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
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TYPE: PRT
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                                                                                                                                             LENGTH: 1488
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   2 WFHVLFYL 9
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RECEPTORS, NUCLEIC ACID MOLECULES ENCODING HUMAN GPCR
PROTEINS, AND USES THEREOF
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57.18;
                                         52.9%;
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71.48;
                                           Score 36;
Pred. No.
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Pred. No. 3.8e+02
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                                Mismatches
                                            DB 6;
5.7e+02;
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Best Local Similarity
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; ORGANISM: Human
US-60-248-505-1121
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                                                                                                                                                                                                                                                                                                                                                                                                         ; ORGANISM: Homo sapiens PCT-US01-01339-5081
                                                                                                                                                                                      PCT-US01-01335-353
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PCT-US01-01339-5081
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SEQ ID NO 5081
LENGTH: 53
TYPE: PRT
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APPLICANT: Human Genome Sciences, Inc.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 1121
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TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
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                                                                                                                                Sequence 353, Application PC/TUS0101335
GENERAL INFORMATION:
APPLICANT: Human Genome Sciences, Inc., et al.
SEQ ID NO 353
LENGTH: 77
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CURRENT FILING DATE: 2001-03-17
NUMBER OF SEQ ID NOS: 10231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Nucleic Acids, Proteins, an FILE REFERENCE: PC006PCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/60/248,505
CURRENT FILING DATE: 2000-11-15
                        NUMBER OF SEQ ID NOS: 879
SOFTWARE: Patentin Ver. 2
                                                               CURRENT APPLICATION NUMBER: PCT/US01/01335
CURRENT FILING DATE: 2001-01-17
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                                                                                              FILE REFERENCE: PC001PCT
                                                                                                                  TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
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24 IWFKIAFFLS 33
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50.0%;
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Pred. No.
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Pred. No. 5.9e+02;
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; ORGANISM: Homo sapiens PCT-US01-01335-353

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Query Match
Best Local Similarity
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PCT-US01-04098A-1850
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Best Local Similarity 42...
6; Conservative
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PRIOR APPLICATION NUMBER: Not Yet 'Assigned PRIOR FILING DATE: 2001-01-30
PRIOR PRIOR DATE: 2000-11-30
PRIOR FILING DATE: 2000-11-30
PRIOR FILING DATE: 2000-10-20
PRIOR FILING DATE: 2000-10-20
PRIOR FILING DATE: 2000-10-20
PRIOR FILING DATE: 2000-09-15
                              SEQ ID NO 187144
LENGTH: 123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 3960
SOFTWARE: Custom
SEQ ID NO 1850
LENGTH: 95
                                                               CURRENT APPLICATION NUMBER: US/09/270,849B
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 195450
SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                       APPLICANT: Swimmer et al.
TITLE OF INVENTION: Insect genome survey devices
FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Homo sapiens
-US01-04098A-1850
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 equence 1850, Application PC/TUS0104098A ENERAL INFORMATION:
ORGANISM: Artificial Sequence
                 TYPE: PRT
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ITLE OF INVENTION: Novel Nucleic Acids and Polypeptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ILE REFERENCE: 21272-029
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42.9%;
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0; Mismatches
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Pred. No.
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le+02;
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US-09-270-849B-186130
Sequence 186130, Application US/09270849B
GENERAL INFORMATION:
APPLICANT: Swimmer et al.
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Best Local Similarity
Thehes 7; Conserv
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; OTHER INFORMATION: Description
US-09-270-849B-187144
Search completed: June Job time: 592 sec
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                                                                                                              Query Match
Best Local Similarity bz...
Thes 5; Conservative
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CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 195450
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 186130
LENGTH: 62
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Insect genome survey devices
                                                                                                                                                                                                                                           ORGANISM: Artificial Sequence FEATURE:
                                                                          1 VWEHVLFY 8
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28 LWVYVLFY 35
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              4, 2001, 12:21:30
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Pred. No.
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq
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length: 2000000000
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Gapop 10.0 , Gapext 0.5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    VLNASVTSPLLE 12
                                                                                                                                                                                                                                                                                                                                                                                                                       /SIDS6/gcgdata/geneseq,
/SIDS6/gcgdata/geneseq/geneseqp/AA2000
  /SIDS6/gcgdata/geneseq/geneseqp/AA1999.DAT:
/SIDS6/gcgdata/geneseq/geneseqp/AA2000.DAT:
                                                                                                                                                                                                                    SIDS6/gcgdata/geneseq/geneseqp/AA1990.DAT:
                                                                 6/gcgdata/geneseq/geneseqp/AA1994_DAT:
6/gcgdata/geneseq/geneseqp/AA1995_DAT:
6/gcgdata/geneseq/geneseqp/AA1996_DAT:
6/gcgdata/geneseq/geneseqp/AA1997_DAT:
                                                                                                                                                                              /qcgdata/g
                                                                                                                                                                                                                                                                                                                                                                              data/geneseq,
                                                                                                                                                                                                  eneseq/geneseqp/AA1991.DAT:
                                                                                                                                                       eneseq/geneseqp/AA1993.DAT:
                                                                                                                                                                                                                                                               /geneseqp/AA1988.DAT:
                                                                                                                                                                                                                                                                                                             /geneseqp/AAJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (without alignments)
6.463 Million cell updates/sec
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                                                                                                                                                                              92.DAT:
                                             .DAT:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

11	10	9	8	7	6	5	4	ω	2	_	Result No.
55	55	55	55	. 55	55	55	55	55	55	55	Score
100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	Query Match I
374	374	374	374	374	374	344	344	344	261	12	Query Match Length DB
19	19	17	17	13	13	21	17	13	20	19	80
W97833	W80448	W00859	R91438	R22550	R20811	Y96183	R91439	R22549	Y33183	W60553	ID
Human Fc receptor	Human Fc receptor	Human FcRI (cDNA c	Human FcRI (cDNA c	Human macrophage-s	Human macrophage-s	Human macrophage-s	Human FcRI (cDNA c	✓ Human macrophage-s	Human sFc-gammaRI	Oligopeptide from	Description

44	43	42	41	40	39	38	37	36	35	34	ü	32	31	30	29	28	27	26	25	24	<u>ي</u>	22	21	20	19	89	17	16	5	14	<u>٦</u>	12
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3 2 2	32	32	32	33	ω ω	ω	3	ω	ω ω	ω ω	ω ω	33.	ω	ω ω	ω ω	34	34	34	34	34	34	3 4	34	34	34	36	36	36	55	55	55	55
58.2 58.2	58.2	8		.0	.0	.0			.0		.0		0			:-	Ξ.	61.8	:	:-	:-	\mathbf{r}	_		:-	5	65.5	•	•		100.0	100.0
176 181	176	139	139	986	788	730	730	377	350	350	144	139	131	131	94	443	442	410	375	374	365	364	322	-187	126	329	329	292	399	374	374	374
21 21	21	21	21	18					15	8	21	21	19	19	21	21	21	12	21	21	21	21	20	19	22	21	21	21	21	21	21	19
G38740 G20313	G20314	G37898	G04555	W13670	W75919	W13669	P80618	W86195	R63143	P70645	G40794	G35305	W37144	W58851	G35306	G47109	G47082	R12428	G47110	G47083	G47111	G47084	W85718	W54052	в63956	Y82015	B28626	B28636	B43683	Y96226	Y96134	W97834 .
																						•					-					
													٠.																			
Arabidopsis Arabidopsis	Arabidopsis	Arabidopsis	Arabidopsis	C-proteinase	<pre>C-proteinase</pre>	<pre>C-proteinase</pre>	Human Bone Morphog	Human Fc receptor	Glycoprotein 63	dorabi	mays	Zea mays protein	Human AC22	Human AC22	Zea mays protein	Arabidopsis	Arabidopsis	Hybrid Fc(gamma)RI	Arabidopsis	Arabidopsis	Arabidopsis	Arabidopsis	el	C. parvum p23	Human prostate can	Human immunogenic	Human B11Ag1	.Human breast	Human cancer	Human high affinit	Human macr	· Human Fc receptor

ALIGNMENTS

PT	ΡŢ	달 :	P	×	DR	XX	ΡI	×	PΑ	×	PR	×	ΡF	×	PD	×	PN	×	SO	X	XW .	ΚW	ΚW	X	DE	×	DT	×	AC .	X	ID	W60553	RESULT
oodies	ide derived from the target, useful for, e.g.		Identifying peptide(s) binding specifically to protein target -		WPI; 1998-240964/21.		De Kruif CA, Logtenberg T;		(UYUT-) RIJKSUNIV UTRECHT.	,	08-OCT-1996; 96EP-0202791.		07-OCT-1997; 97WO-NL00557.		16-APR-1998		WO9815833-A1.		Synthetic.			_	Extracellular domain; CD64; identification; antibody;		Oligopeptide from extracellular domain of CD64.		18-AUG-1998 (first entry) '		· W60553;		W60553 standard; peptide; 12 AA.	53	LT 1

method for the identification of a peptide able to bind specifically to a target protein. The method comprises displaying the peptide on the surface of a replicable display package, synthesising oligopeptides derived from the target protein on a solid phase, and testing for binding between the peptide and oligopeptides. The method is used to screen large peptide libraries, especially to detect antibodies, or their fragments, that bind to cell markers or thousand sifferent forms of the same protein, including bispecific antibodies that bind to two non-overlapping epitopes on the same monomeric antigen or two epitopes on different molecules. The genes/foligonucleotides that encode

Synthetic oligopeptides W60537-61 are derived from the extracellular domain of CD64. They were synthesised on 25 polyethylene rods as adjacent, non-overlapping sets. Peptide specific phages were isolate.

The specification describes a

Example 1; Page 29; 40pp;

English

affinity binding to the CD64-covered rods

Query Match

Length 12;

Sequence

12

ΑĀ,

by mutation in (anti)oncogenes. antibodies. Selected antibodies

antibodies. Selected antibodies are used, e.g. for immunohistochemical or immunofluorescent analysis, and also to detect cell transformation caused

peptides can be isolated and used for recombinant production of

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
Three-dimensional structures and models of Fc receptors, computer-assisted drug design
                                                                                   Baell JB, I
                                           WPI; 1999-539978/45
                                                                                                                                                         11-SEP-1998;
06-FEB-1998;
                                                                                                                                                                                                     04-FEB-1999;
                                                                                                                                                                                                                                                           W09940117-A1
                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                  anti-inflammatory;
                                                                                                                                                                                                                                                                                                                             tissue damage; hypersensitivity; inflammatory cell recruitment; inflammatory modulator; Fc-gammaRIa; immune function regulation;
                                                                                                                                                                                                                                                                                                                                                          atomic coordinate; bioactive compound design; computer-assisted; drug design; therapy; inhibitor; Fc-gammaR; Fc-epsilonR; IgG; Ig
                                                                                                                                                                                                                                                                                                                                                                                                                  Human sFc-gammaRI protein fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                             15~NOV-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Y33183 standard; Protein;
                                                                                                                           (ILEX-') ILEXUS
                                                                                                                                                                                                                                12-AUG-1999
                                                                                                                                                                                                                                                                                                                                                                                      FC receptor; FC-gammaRIIa; human; FCR; model; three-dimension; 3-D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
nes 12; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 VLNASVTSPLLE 12
|||||||||||
| vlnasvtsplle 12
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                                                                                   Epa V, Garret, McCarthy TD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                              PTY LTD .-
                                                                                                                                                         98US-0099994
98US-0073972
                                                                                                                                                                                                    99WO-IB00367.
                                                                                                Garrett TPJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     261
                                                                                   TPJ, Hogarth PM, Matthews McKenzie IFC, Pietersz GA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 55; DB 19;
Pred. No. 6.2e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                Matthews
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
              useful
                                                                                                                                                                                                                                                                                                                                                          IgG; IgE;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
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This amino acid sequence was predicted from the cDNA sequence of CDNA clone p98/X2. It differs from the sequence predicted from clone p185 (see Q211.78) at position 25, p135 encodes a Ser residue while clone p98/X2 predicts a Thr residue. At position 58, p90

Ser residue 58, p90

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    R22549
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This invention describes a novel model of an FC receptor (FCR) protein CC representing a three-dimensional (3-D) structure that substantially CC conforms to the specified atomic coordinates. Computer model images of the FCR can be used to design bioactive chemical compounds, e.g. cligonucleotides, peptides, peptidomimetics and small organic molecules, by computer-assisted methods of drug design. Therapeutic compositions CC by computer-assisted methods of drug design. Therapeutic compositions CC that inhibit the activity of FC-qammaR or FC-epsilonR can be used to reduce IgG- or IgE-mediated tissue damage, e.g. IgG- or IgE-mediated hypersensitivity, recruitment of inflammatory cells or release of CC inflammatory modulators. The therapeutic compositions can also be used to treat disease in animals. The knowledge of the 3-D structure of FC-gamma CC RIA and models of other FCR provides a means for designing and producing CC compounds that regulate immune function and inflammation in an animal, CC including humans (i.e. structure based drug design). For example, including can be designed to block binding of immunoglobulin to the processor of the structure based drug design and models.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local S
Matches 12
                                                                                                              New CD53 cell surface antigen and DNA encoding immuno-therapy and diagnosis of haematopoietic
                                                                                                                                                                  WPI; 1992-056864/07
N-PSDB; Q21179.
                                                                                                                                                                                                                      Seed
                                                                                                                                                                                                                                                                                        13-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         an Fc receptor protein using various computer programs and models. The products of the invention have anti-inflammatory and immunoprotective activity. This sequence represents the human sFC-gammaRI protein fragment described in the method of the invention.
                                                                                Example
                                                                                                                                                                                                                                                      (GEHO-) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                        15-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                            23-JAN-1992
                                                                                                                                                                                                                                                                                                                                                                                            WO9201049-A
                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rapid immunoselection cloning technique; cell surface antigen; immunodiagnosis; high affinity receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human macrophage-specific FcRI receptor encoded by clone p98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  R22549;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    R22549 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 11; Page 316-317; 326pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    174 vlnasvtsplle 185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 VLNASVTSPLLE
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                                                                                                                                                                                                                    В,
                                                                                10;
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                                                                                                                                                                                                                    Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            261
                                                                            Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; ilarity 100.0%; Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                          90US-0553759
                                                                                                                                                                                                                                                                                                                          90WO-US04986
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12
                                                                                                                                                                                                                      Amiot M;
                                                                            160pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    344
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 55; DB 20;
Pred. No. 0.0036;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                neoplasms,
                                                                                                                                it - for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 261;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                  etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
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Query Match
Best Local Similarity
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The amino acid sequence (R91439) of human FCRI was detd. from a cDNA clone, p98 (T14718), obtd. from a cDNA library using an cDNA clone, p98 (T14718), obtd. from a cDNA library using an common control clothing method. FCRI is a high affinity receptor for the FC portion of IgG, normally located on cell surfaces of macrophages. 2 Other cDNA clones, p135 (T14717) and p90 (T14719), coded for a variants of the FCRI sequence; the C-terminal sequence of the p98 product is truncated compared with those of the p135
                                                                                                                                                                                                                Cloning of cDNA encoding cell surface antigen of diagnostic and therapeutic proteins
                                                                                                                                                                                                                                                                                  N-PSDB; T14718.
                                                                                                                                                                                                                                                                                                          WPI:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-DEC-1992;
25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane-spanning domain, and a short, highly charged cytoplasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and six
                                                                                                                                                                                                                                                                                                                                               Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                09-APR-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US5506126-A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R91439;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R91439 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cys residues distributed among three C2 set Ig-related domains: A fusion protein of FcRI and a receptor ligand will be helpful to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (see Q21180) predicts Val and p98/X2 predicts Leu. See predicted from all 3 clones show the typical features
                                                                                                                                                                         Example 10; Column 55-56; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                13-JUL-1989
13-JUL-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cell surface antigen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                      (GEHO ) GEN HOSPITAL
                                                                                                                                                                                                                                                                                                                                                                                                                               18-OCT-1993
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                                                                                                                                                                                                                                                                                                     1996-200279/20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FCRI (cDNA clone p98 product).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       the potency of antibodies
                                                                                                                                                                                                                                                                                                                                               Seed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    344 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                             89US-0379076.
90US-0553759.
93US-0139273.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         88US-0160416
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                                                                                                                                                                                                                                                                                                                                               В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note=
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ector; FcRI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            amino acid 58 is Leu in clone pl35 translated product"
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  immunoselection; immunotherapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              receptor.
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                                                                                                                                                                                                                                  useful for isolation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 344;
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ses of a type I
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B 8

Misc-difference

/note= 213

"encoded

by AAT"

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33
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Y96183
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Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              and p90 products (see also R91438 and W00859). if of FCRI and a receptor ligand will be helpful in potency of antibodies in therapy.
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                                                                                                                                                                                                                                                                                                                                                                                                                              amyloidosis; multiple sclerosis; parasitic disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                       panning; immunodiagnosis; diagnosis; immunotherapy; gene ty
immune disorder; infection; asthma; immune-complex disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human macrophage-specific FcRI.
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nilarity 100.0%;
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= 60
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/note-
140
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                                                                                                               /note=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            cell surface antigen;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 344;
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                                                                                                                                                                                                                                                                                                                                                                                                                               receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                 therapy;
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Misc-difference

/note= 216

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sclerosis. FCRI is a high affinity receptor for the Fc portion of IgG, normally located on the cell surfaces of macrophages. The ability to interfere with such bonding, or to cause it to occur on surfaces other than macrophages, is useful in therapy. A fusion protein of FCRI and a receptor ligand will be helpful to increase
                             surfaces other than macrophages, is useful protein of FCRI and a receptor ligand will the potencies of antibodies in therapy.
                                                                                                                                                                                                                   (panning on) an antibody-coated substrate such as a culture dish. CSA nucleic acids isolated by the method of the invention, and the proteins they encode, are useful for immunodiagnostic and
                                                                                                                                                                                                                                                                                                                             The present sequence is that of a human macrophage specific FCRI, as deduced from cDNA clone p135 (see A50631), which was isolated from a cDNA library expressed in COS cells using a novel method of
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Isolated nucleic acid molecule encoding the CD19 cell surface antigen, useful for immunodiagnosis and immunotherapy of immune-mediated infections or disorders, e.g. asthma, immune-complex disease, parasiti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Stamenkovic I,
                                                                                                                                                                                immunotherapeutic applications, including the diagnosis and disorders in treatment of immune-mediated infections, diseases, and disorders in
                                                                                                                                                                                                                                                                          physical selection of cells expressing the antigen by adhesion to
                                                                                                                                                                                                                                                                                            the invention designed to isolate CSA nucleic acids. The met
based upon transient expression of a CSA in eukaryotic cells
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23-MAR-1990,
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                                                                                                                                           nimals, including humans. These disorders include asthma, mmune-complex disease, amyloidosis, parasitic diseases or multiple
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5-FEB-1988
                                                                                                                                                                including humans.
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90US-0498809
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338
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333
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305
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268
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220
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                                                                                                                                                                                                                                                                                                                                                                                                            English.
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                                                                                                                                                                                                                                                                                                                  The method is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   parasitic
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Query Match 100.0%; Best Local Similarity 100.0%; Matches 12; Conservative (

0

Mismatches

Score 55; DB 21; Pred. No. 0.0052;

DB 21;

Length 344; Indels

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Gaps

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1 VLNASVTSPLLE 12

Sequence

344 AA;

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RESULT
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R20811
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Best Local Similarity 100
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             position 25; p135 encodes a Ser residue and the other two clones predicts a Thr residue. At position 58, p135 predicts Leu and p90 predicts Val. Sequences predicted from all 3 clones show the typical features of a type I integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane-spanning domain, and a short, highly charged cytoplasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and six Cys residues distributed among three C2 set Ig-related domains. A fusion protein of FCRI and a receptor ligand will be helpful to increase the potency of antibodies in therapy.
                          21-MAY-1992 '(first entry)
                                                                                                        R22550;
                                                                                                                                                                        R22550 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cDNA clone plas. It differs from the sequence predicted from clones p99/X2 and p90 (see Q21179 and Q21180, respectively) at position 25; plas encodes a Ser residence and the position 25; plas encodes a Ser residence and the position 25; plas encodes a Ser residence and the position 25; plas encodes a Ser residence and the position 25; plas encodes a Ser residence and the position 25; plas encodes a Ser residence and the position 25; plas encodes a Ser residence and the position 25; plas encodes a Ser residence and the position 25; plas encodes a Ser residence and the position 25; plas encodes a Ser residence and the position 25; plas encodes a Ser residence and the position 25; plas encodes a Ser residence and the position 25; plas encodes a Ser residence and the position 25; plas encodes a Ser residence and the position 25; plas encodes a Ser residence and the position 25; plas encodes a Ser residence and the position 25; plas encodes a Ser residence and the position 25; plas encodes a Ser residence and the position 25; plas encodes a Ser residence and the position 25; plas encodes a Ser residence and the position 25; plas encodes a Ser residence and the position 25; plas encodes a Ser residence and the position 25; plas encodes and the positi
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         R20811 standard; Protein; 374 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New CD53 cell surface antigen and DNA encoding it - for immuno-therapy and diagnosis of haematopoietic neoplasms,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1992-056864/07
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rapid immunoselection cloning technique; cell surface antigen; immunodlagnosis; high affinity receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human macrophage-specific FcRI receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-MAY-1992
                                                                                                                                                                                                                                                                                                                                                           193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     193 vlnasvtsplle
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     374 AA;
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                                                                                                                                                                           Protein; 374
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 55; DB 13; -Pred. No. 0.0058;
                                                                                                                                                                           A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 374;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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Human

macrophage-specific FcRI receptor

encoded by clone p90

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RESULT
R91438
                                                                                                                                                                                                                                                                                                                                                                                                                      Š
 ETT FTT KW
                                                                                                                                                                                                                                                                                                                                                                                       В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 12; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This amino acid sequence was predicted from the cDNA sequence of cDNA clone p90. It differs from the sequence predicted from clone p135 (see 021178) at position 25; p135 encodes a Ser residue and clone p90 predicts a Thr residue. At position 58, p90 predicts val while clones p135 and p98/X2 (see 021179) both predict Leu. Sequences predicted from all 3 clones show the typical features of a type I integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane-spanning domain, and a short, highly charged cytoplasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and six Cys residues distributed among three C2 set Ig-related domains. A fusion protein of FGRI and a receptor ligand will be believed beliefil to increase the potency of antibodies in the server.
                                                                                                                                  Cell surface antigen; cloning; immunoselection; immunotherapy; therapy; diagnosis; vector; FcRI; Fc receptor.
                                                                                                                                                                                           Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
Misc-difference
                                                                                                     Homo sapiens
                                                                                                                                                                                                                          30-OCT-1996 (first entry)
                                                                                                                                                                                                                                                             R91438
                                                                                                                                                                                                                                                                                                R91438 standard; Protein; 374 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New CD53 cell surface antigen and DNA encoding it - for immuno-therapy and diagnosis of haematopoietic neoplasms,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           N-PSDB; Q21180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Seed B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-JÜL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rapid immunoselection cloning technique; cell surface antigen; immunodiagnosis; high affinity receptor.
                                                 Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  helpfül to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 10; Page 94a; 160pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1992-056864/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-JUL-1990; · 90US-0553759
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          23-JAN-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           W09201049-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (GEHO-) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                       193
                                                                                                                                                                                                                                                                                                                                                                                                       1 VLNASVTSPLLE 12
                                                                                                                                                                                                                                                                                                                                   8
                                                                                                                                                                                                                                                                                                                                                                                     vlnasvtsplie 204
                                                                                                                                                                                       FCRI (cDNA clone p135 product).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  increase the potency of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      90WO-US04986
                                               Location/Qualifiers
                               /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Amiot M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100
            "amino acid 25 is Thr in clone p90 and p98 translated products"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           .0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 55; DB Pred. No. 0.0 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 13;
0.0058;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 374;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
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В
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                                                                                                                                                                                                                                                                                                                                                                                                  SXCCCCCCCXXXXIX
                                                                                                                                                                                           DXXX
                                                                                                                                                                                                                                                  W00859
                                                                                                                                                                                                                                                              RESULT
                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
Matches 12; Conser
 01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13-JUL-1989;
13-JUL-1990;
18-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                    The amino acid sequence (R91438) of human FCRI was detd. from a cDNA clone, p135 (T14717), obtd. from a cDNA library using an immunoselection cloning method. FCRI is a high affinity receptor for the FC portion of IgG, normally located on cell surfaces of macrophages. Another isolated cDNA clone, p90 (T14719), coded for a variant (W00859) of the FCRI sequence, and a third clone, p96 (T14718), coded for an FCRI (R91439) having a different C-terminal sequence. A fusion protein of FCRI and a receptor ligand will be helpful in increasing the potency of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cloning of cDNA encoding cell surface antigen of diagnostic and therapeutic proteins
                                            25-FEB-1988
                                                                 09-APR-1996
                                                                                        US5506126-A
                                                                                                             Homo sapiens
                                                                                                                                  therapy; diagnosis;
                                                                                                                                              Cell surface antigen; cloning; immunoselection; immunotherapy;
                                                                                                                                                                   Human FcRI (cDNA clone p90 product).
                                                                                                                                                                                           30-OCT-1996
                                                                                                                                                                                                                 W00859;
                                                                                                                                                                                                                                      W00859 standard; Protein; 374
                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 10; Column 55-56; 79pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             N-PSDB; T14717.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1996-200279/20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-DEC-1992;
25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   09-APR-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US5506126-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (GEHO ) GEN HOSPITAL CORP.
                                                                                                                                                                                                                                                                                              193 vlnasvtsplle 204
                                                                                                                                                                                                                                                                                                        1 VLNASVTSPLLE 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Seed B;
                                                                                                                                                                                                                                                                                                                                                                                                  374
                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                           (first entry)
92US-0983647.
88US-0160416.
89US-0379076.
                                                                                                                                                                                                                                                                                                                                                                                                  AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          88US-0160416.
89US-0379076.
90US-0553759.
93US-0139273.
                                            88US-0160416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        92US-0983647
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             88US-0160416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              vector; FcRI; Fc receptor
                                                                                                                                                                                                                                                                                                                                                    100.0%;
                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                     Score 55;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     English.
                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                   0.0058;
                                                                                                                                                                                                                                                                                                                                                               DB 17;
                                                                                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        useful for isolation
                                                                                                                                                                                                                                                                                                                                                               Length
                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                 374;
                                                                                                                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                                                                                                                                                        98d
                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                           0;
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RESULT
W80448
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
Matches 12; Conser
Misc-difference
                          Misc-difference
                                                    Misc-difference
                                                                                Misc-difference
                                                                                                           Misc-difference
                                                                                                                                                              Misc-difference
                                                                                                                                                                                                                                             Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The amino acid sequence (W00859) of human FCRI was detd. from a CDNA clone, p90 (T14719), obtd. from a cDNA library using an immunoselection cloning method. FCRI is a high affinity receptor for the FC portion of IgG, normally located on cell surfaces of macrophages. Other cDNA clones (see also T14717-18) coded for variants (R91438-39) of the FCRI sequence. A fusion protein of FCRI and a receptor ligand will be helpful in increasing the potency of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cloning of cDNA encoding cell of diagnostic and therapeutic
                                                                                                                                   Misc-difference
                                                                                                                                                                                        Misc-difference
                                                                                                                                                                                                                  Misc-difference 44
                                                                                                                                                                                                                                                                       Misc-difference
                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                          cloning
                                                                                                                                                                                                                                                                                                                                                      Fc receptor I; FcRI;
                                                                                                                                                                                                                                                                                                                                                                                Human Fc receptor I.
                                                                                                                                                                                                                                                                                                                                                                                                            07-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                       W80448;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               W80448 standard; Protein; 374 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 10; Column 55-56; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1996-200279/20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-JUL-1990;
18-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 193 vlnasvtsplle 204
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 VLNASVTSPLLE 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                     /note-
85
                          /note=
141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    90US-0553759.
93US-0139273.
/note=
159
                                                    /note=
103
                                                                                /note=
99
                                                                                                                                                               60
                                                                                                                                                                                          45
                                                                                                                                                                                                                                             /note=
23
                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                               'note-
                                                                                                                                                                            'note=
                                                                                                                                                                                                     'note='
                                                                                                                                                                                                                              note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%;
                                                                                                                                                                                                                                                                                                                                               cell surface antigen; lymphocyte; human;
            "encoded
                                                                                                                                                                                                                                                         "encoded by TGG"
                                                                "encoded
                                                                                                                  "encoded by
                                      "encoded by
                                                                                                                                                                                                                              "encoded by CTC"
                                                                                                                                              "encoded by CCC"
                                                                                                                                                                         "encoded by CTG"
                                                                                                                                                                                                    "encoded by GAC"
                                                                                          "encoded by TCC"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 55; DB 17;
Pred. No. 0.0058;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            surface antigen - useful for isolation
            bу
                                                               ьy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                       CCC"
                                                                CAA"
                                                                                                                      AAT"
              GGC"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 374;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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RESULT -W97833 AX I

W97833;

W97833 standard;

Protein; 374

DЬ Qy

193 vlnasvtsplle 204

1 VLNASVTSPLLE 12

Matches

12;

Query Match Best Local Similarity

100.0%; ilarity 100.0%; Conservative

0

Mismatches

0;

0; Gaps

0;

Score 55; DB 19; Pred. No. 0.0058;

Length Indels

374;

Sequence

374

AA;

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33333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               XΥ
                                                                                    the isolation and cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V63442-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated genes can be expressed in a prokaryotic or eukaryotic host cells to produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V63444) which allow the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13-JUL-1989;
13-JUL-1990;
21-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially human lymphocyte antigenic sequences
                                                                                                                                                                                                                                                            expression libraries is based on transient expression of an antigen by in eukaryotic cells and selection of cells expressing the antigen by adhesion to an antibody-coated substrate. The method is useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Misc-difference
                                                   immunotherapeutic applications, treatment of immune-mediated in
                                                                                                                                                                                                                                                                                                                     induction therapy. were also obtained.
                                                                                                                                                                                                                                                                                                                                                                                                                This is the amino acid sequence of human Fc receptor I (FcRI), deduced from cDNA clone pl35 (see V63456) isolated using a rapi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-DEC-1992;
25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         03-NOV-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US5830731-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Misc-difference
                                                                                                                                                                                                                                                                                                                 from a single patient undergoing extracorporeal interleukin-2 induction therapy. Clones p90 (see x07372) and p98 (see x07373) were also obtained. A novel method for cloning CDNAs from mamma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 10; Column 53-54; 75pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   N-PSDB; V63456
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1998-609251/51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                            immunoselection cloning method from a cDNA library expressed in opening.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (GEHO ) GEN HOSPITAL CORP
                                 including
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Seed B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   90US-0553759
97US-0861205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            92US-0983647
88US-0160416
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256
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176
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                                 humans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "encoded by GGG"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "encoded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "encoded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "encoded
                                                       infections
                                                                    including the diagnosis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     bу
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GTC"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATG"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAC"
                                                                                                                                                                                                                                                                                                                     from mammalian
                                                       disorders
                                                                                                                                                                                                                                                                                                                                                                                                                (FCRI), as
g a rapid
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in cos

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25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
                                                                 This is the amino acid sequence of human Fc receptor I (FcRI), deduced from cDNA clone p90 (see X07372) isolated using a rapi immunoselection cloning method from a cDNA library expressed i
                                                                                                                                                New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially human lymphocyte antigenic sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note-
Misc-difference 103
                                                                                                                                                                                                       WPI; 1998-609251/51.
N-PSDB; X07372.
                                                                                                                                                                                                                                                                                                                                                        01-DEC-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                03-NOV-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                          US5830731-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cloning.
                                                                                                                      Example 10; Column 53-54; 75pp; English.
                                                                                                                                                                                                                                              Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                     21-MAY-1997;
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              were also obtained
                                                                                                                                                                                                                                                                        (GEHO ) GEN HOSPITAL
                                                                                                                                                                                                                                                                                                    21-MAY-1997;
                          induction therapy
                                          from a single patient undergoing extracorporeal
                                                     The cDNA library was constructed from polyA RNA of cells
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88US-0160416.
89US-0379076.
90US-0553759.
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. 77
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176
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                            Clones p135
                                                                                                                                                                                                                                                                         CORP.
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based on transient expression
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              (see V63456) and p98 (see X07373) hod for cloning cDNAs from mammal
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Misc-difference 141
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes see V63442-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated genes can be expressed in a prokaryotic or eukaryotic host cells to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                in eukaryotic cells and selection of cells expressing adhesion to an antibody-coated substrate. The method the isolation and cloning of any protein which can be
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   expression libraries is based on transient expression of an antigen in eukaryotic cells and selection of cells expressing the antigen by adhesion to an antibody-coated substrate. The method is useful for the isolation and cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V63442-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated genes can be expressed in a prokaryotic or eukaryotic host cells to produce the encoded protein. The invention also provides high cefficiency expression vectors (see V63441 and V63444) which allow the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and immunother apeutic applications, including the diagnosts and
                                                                                                                                                                                                                                                                                                                                                                                             Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 12; Conservative
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13-JUL-1990;
21-MAY-1997;
                Macrophage; FCRI; cell surface antigen; human; immunoselection; panning; immunodiagnosis; diagnosis; immunotherapy; gene therapy; immune disorder; infection; asthma; immune-complex disease;
  amyloidosis;
                                                                                                Human macrophage-specific FcRI
                                                                                                                                         19-DEC-2000
                                                                                                                                                                                                                   Y96134 standard; Protein; 374 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This is the amino acid sequence of human Fc receptor I (FCRI) deduced from cDNA clone p98/X2 (see X07373) isolated using a immunoselection cloning method from a cDNA library expressed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially human lymphocyte antigenic sequences
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25-FEB-1988;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cells. The cDNA library was constructed from polyA RNA of cells from a single patient undergoing extracorporeal interleukin-2 induction therapy. Clones p135 (see V63456) and p90 (see X07372) were also obtained. A novel method for cloning cDNAs from mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 10; Column 53-54; 75pp; English.
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N-PSDB; X07373.
                                                                                                                                                                                                                                                                                                                       193 vlnasvtsplle 204
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multiple sclerosis;
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88US-0160416.
89US-0379076.
90US-0553759.
97US-0861205.
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  parasitic disease;
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  Fc receptor
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01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
23-MAR-1990;
13-JUL-1990;
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Misc-difference 122
       Stamenkovic I,
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Misc-difference 126
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Misc-difference 123
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Misc-difference 116
                     (GEHO ) GEN HOSPITAL CORP
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                                                                                                                                                                                                                                                                                                                                                                                                  CAA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                TCC"
                                                                                                                                                                                                                                                                                                                                                                                                                                                               CAC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                ccc"
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WPI; 2000-586382/55 N-PSDB; A50592.

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RESULT 14
Y96226
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 12; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     immune-complex disease, amyloidosis, parasitic diseases or multiple sclerosis. FcRI is a high affinity receptor for the Fc portion of IgG, normally located on the cell surfaces of macrophages. The ability to interfere with such bonding, or to cause it to occur on surfaces other than macrophages, is useful in therapy. A fusion protein of FcRI and a receptor ligand will be helpful to increase
                                                                                                              07-JUN-2000
                                                                                                                                                                                         Human; high affinity Fc receptor; FcgammaRI; immunoglobulin; infection; immune response; CD64; monocyte; macrophage; neutrophil; eosinophil; HIV; IGG; immunosuppressive; antirheumatic; cytostatic; antiinflammatory; AIDS; SLE; multiple myeloma; rheumatoid arthritis;
                                                                                                                                                                                                                                                                         11-SEP-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                  193
         N-PSDB; A27466
                                                                03-DEC-1998;
                                                                                      03-DEC-1998;
                                                                                                                                     EP1006183-A1
                                                                                                                                                                                systemic lupus
                                                                                                                                                                                                                                                                                                                        Y96226 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                    WPI; 2000-367968/32
                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                  Human high affinity Fc receptor, FcgammaRI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           the potencies of antibodies in therapy.
                                          (PLAC ) MAX PLANCK GES
                                                                                                                                                                                                                                                                                                                                                                                                      1 VLNASVTSPLLE 12
                                                                                                                                                                                                                                                                                                                                                                                vlnasvtsplle 204
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                  98EP-0122969
                                                                                                                                                                                erythematosus; tumour.
                                                                                       98EP-0122969
                                                                                                                                                                                                                                                                                                                       Protein; 374 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%;
                                           FOERDERUNG WISSENSCHAFTEN
                                                                                                                                                                                                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 55;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                        0.0058;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                   21;
                                                                                                                                                                                                                                                                                                                                                                                                                              0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 374;
                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                             0;
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may be modified to produce recombinant versions. The recombinant Ro receptor consist only of the extracellular portion of the receptor and are not glycosylated i.e. they do not have transmembrane domains or signal peptides. The recombinant proteins may be used in immunoassays t determine the immune status of patients with chronic diseases of the immune system, e.g. AIDS, systemic lupus erythematosus (SLE), multiple myeloma (MM), or rheumatoid arthritis. In addition, pharmaceutical compositions containing recombinant proteins may be not the treat or compositions containing recombinant proteins may be not the treat or compositions containing recombinant proteins may be not the treat or compositions containing recombinant proteins may be not the treat or compositions containing recombinant proteins may be not the treat or compositions containing the proteins may be not the treat or compositions containing the proteins may be not the proteins and the proteins are proteins and the proteins and the proteins are proteins are proteins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cells expressing Fc receptors. FcgammaRI molecules are expressed by monocytes and macrophages, but expression can also be induced on neutrophils and eosinophils. Upon Fc receptor activation, immune effector pathways are activated, leading to immune response. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       glycosylation, cancer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel Fc receptor lacking transmembrane domains, a signal peptide, and glycosylation, useful for diagnosing and treating immune disorders and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure;
                                                                                    rheumatoid arthritis
                                                                                                                     compositions containing recombinant proteins may be used to treat or prevent autoimmune diseases, allergies or tumours, especially AIDS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  role in defending the body against infections. First, pathogens are opsonised by serum immunoglobulins. The resulting complex then binds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present sequence is the human high affinity Fc receptor, FcgammaRI. FcgammaRI is also known as CD64. Fc receptors play an important
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page 26-28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              60pp; English
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physical selection of cells expressing the antigen by adhesion to (panning on) an antibody-coated substrate such as a culture dish. CSA nucleic acids isolated by the method of the invention, and the proteins they encode, are useful for immunodiagnostic and immunotherapeutic applications, including the diagnosis and

animals,

treatment of immune-mediated infections,

including humans.

These disorders include asthma,

diseases, and disorders in

multiple ortion of

Sequence

The present sequence is that of a human macrophage specific FCRI, as deduced from cDNA clone pl35 (see A50592), which was isolated from a cDNA library expressed in COS cells using a novel method o

a novel method of

The method is

the invention designed to isolate CSA nucleic acids.

upon transient expression of a CSA in eukaryotic cells and

Example 10; Column 53-55; 75pp; English.

Isolated nucleic acid molecule encoding the CD19 cell surface antigen, useful for immunodiagnosis and immunotherapy of immune-mediated infections or disorders, e.g. asthma, immune-complex disease, parasitic

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RESULT
                                                                                                                        Query Match
Best Local Similarity
Matches 12; Conserv
08-FEB-2001
                   B43683;
                                     B43683 standard; Protein; 399
                                                                                  15
                                                                                                       1 VLNASVTSPLLE 12
                                                                                                                          Conservative
(first entry)
                                                                                                                                  100.0%;
                                                                                                                                           .0%;
                                                                                                                        Score 55; DB 21; I
Pred. No. 0.0058;
Pred. No. 0.0058;
                                                                                                                          0;
                                      Ā
                                                                                                                                           Length 374;
                                                                                                                          Indels
                                                                                                                          0;
                                                                                                                          Gaps
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В

Human cancer associated protein sequence SEQ ID NO:1128 cancer associated gene; cancer antigen; detection; cytostatic; neuroprotective; thrombolytic; proliferative; vulnerary; coagulant; immunomodulator; cancer;

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vasotropic; antipsoriatic; antianglogenic; gene therapy; inflammation; immune disorder; haematopoletic cell disorder; autoimmune disorder; altergic reaction; graft versus host disease; organ rejection; haemostatic; thrombolytic; cardiovascular disorder; infection; neurological disease; drug screening.
WPI; 2000-587533/55
                                                                                                                                                                                                                                                               08-MAR-2000; 2000WO-US05882
                                                                                                                                                                                                                                                                                                                                                                                            WO200055350-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antidiabetic; antiasthmatic; antirheumatic; antiarthritic; antiviral;
antiinflammatory; antithyroid; antiallergic; antibacterial; cardiant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         diagnosis;
                                                                Rosen CA,
                                                                                                                          (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                              12-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          dermatological;
                                                                Ruben
                                                                                                                                                                                              99US-0124270.
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cc antityroid; antialization, antibacterial; antiviral; dermatological; cc neuroprotective; cardiant; thrombolytic; coagulant; nootropic; antipsoriatic and antianglogenic. The polynucleotides and conditions and diagnosing pathological conditions. Polynucleotides, conditions and personable and antagonists from the present cc invention may be used to treat immune disorders by activating or inhibiting the proliferation, differentiation or mobilisation of immune cells, to treat disorders of haematopoietic cells, autoimmune disorders, callergic reactions, graft versus host disease and organ rejection, comodulate haemostatic or thrombolytic activity, modulate inflammation, cc cancers, cardiovascular disorders, neurological disease and bacterial or viral infections. The peptides, nucleotides, authordies, agonists and cantagonists may be also be used in drug screens. C78449 to C78457 and convention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C77607 to C78448 encode the human cancer associated proteins given in B43398 to B44239. The proteins can have activities based on the tissues and cells the genes are expressed in Example of activities include: cytostatic; proliferative; vulnerary; immunomodulator; antidiabetic; antiasthmatic; antirheumatic; antiarthritic; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel isolated nucleic acids comprising sequences encoding peptides useful for treating or diagnosing e.g. cancer \cdot
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 11; Page 1739-1740; 2352pp; English.
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Search completed: June 4, Job time: 199 sec 망 Š Ouery Match 100.0%; Score 55; DB 21; Length 399; Best Local Similarity 100.0%; Pred. No. 0.0063; Matches 12; Conservative 0; Mismatches 0; Indels 1 VLNASVTSPLLE 12 |||||||||| | 218 vlnasvtsplle:229 2001, 12:13:26 0; Gaps

0;

Sequence

399 AA;

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Result
No.
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Maximum DB
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Title:
Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Run on:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             rotal number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              seq
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length: 2000000000
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55
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             185757 seqs, 19210857 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd
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/cgn2_6/ptodata/2/iaa/PCTUS_COMB.pep:*
/cgn2_6/ptodata/2/iaa/backfiles1.pep:*
                                                                                                                                                                                                                                                                                                                                                                                                                                      Length
                                                                                               1362
20
193
                                                                                                                                                                                                                                                                                                                                                                                                                                        DB
US-08-572-225-1
US-08-785-164-15
US-08-785-164-6
US-08-785-164-6
US-08-785-164-6
US-08-785-164-6
US-08-785-164-14
US-08-785-164-14
US-08-785-164-14
US-08-785-164-16
                                                                                                                                                                                                                                                                                                                                                                       US-08-332-562A-9
US-08-764-100-27
US-08-721-488-8
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Sequence
                                                                                                                                                                                                                                                                                                                                                          Patent
                                                                                                                                                                                                                                                                                                                                                                                                                                     Description
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                                 Sequence
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                                                                                                                                                                                                                                                                                                                                                          No.
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12, Appl
6, Appli
6, Appli
e 1, Appli
e 1, Appli
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5, Appl
5, Appl
3, Appl
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	56.4	56.4	56.4	56.4	56.4	56.4	56.4	56.4	56.4	56.4	56.4	56.4	56.4	56.4	56.4	56.4	56.4	56.4	
	617	617	617	617	617	572	572	535	390	382	382	382	382	285	247	247	247	194	
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ALIGNMENTS	US-08-279-700-10	US-08-279-700-8	US-08-279-700-6	US-08-279-700-4	US-08-279-700-2	PCT-US91-08177-19	PCT-US91-08177-11	US-08-564-972-1	US-09-036-987A-8	PCT-US96-01444-7	US-08-555-268A-14	US-08-894-236-7	US-08-415-818-7	US-09-141-821-4	US-08-904-686A-44	US-08-384-616-44	US-08-324-977-44	US-08-466-604-29	
	Sequence 10, Appl	Sequence 8, Appli	Sequence 6, Appli	Sequence 4, Appli	Sequence 2, Appli	Sequence 19, Appl	Sequence 11, Appl	Sequence 1, Appli	Sequence 8, Appli	Sequence 7, Appli	Sequence 14, Appl	Sequence 7, Appli	Sequence 7, Appli	Sequence 4, Appli	Sequence 44, Appl	Sequence 44, Appl	Sequence 44, Appl	Sequence 29, Appl	

RESULT

US-08-332-562A-94

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TOPOLOGY:
US-08-332-562A-94
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 9. Patent No.
                                                                                                  INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                CLASSIFICALION.

PRIOR APPLICATION NUMBER: US 07/896,457

APPLICATION TOMBER: US 07/896,457

TITME DATE: 27-MAY-1992
                                                                                                                            FILING DATE: 27-MAY-1992
ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 54270/119/GRHA
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
TELEPAX: (202)672-5399
                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: HULETT, MARK D. TITLE OF INVENTION: FC RECEPTOR FOR IMMUNOGLOBULIN
                                           SEQUENCE CHARACTERISTICS:
LENGTH: 91 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
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                                                                                                     TELEFAX: (20-, 75)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 3000 K S
CITY: Washington
STATE: D.C.
                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                STRANDEDNESS:
                                                                                                                                                                                                                                                                                                 FILING DATE: 31 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    94,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                'RY: USA
20007-5109
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5985599
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HOGARTH, Mark P.
HIBBS, Margaret L.
                                                                                                                                                                                                                                                                                                                                                              PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BONADONNA, Lisa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MCKENZIE,
               linear
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N: 435
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RESULT 3
US-08-721-488-8
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Sequence 8, Application US/08721488
                                                                                                                                                         Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity 66.7%; Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 17-MAR-1993
APPLICATION NUMBER: GB 9206016.9
FILING DATE: 19-MAR-1992
ATTORNEY/AGENT INFORMATION:
NAME: NO 5773700ris, Allen E:
REGISTRATION NUMBER: 34.490
REGISTRATION NUMBER: 34.490
                                                                                                                                                                                                                                                                                                                                 TELEFAX: (415) 857-1125
NFORMATION FOR SEQ ID NO: 27:
                                                                             ||| ||:||
2230 LNALAASPILE 2240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION: TELEPHONE: (415) 354-3592
                                                                                                                                                                                                                                      MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MBER OF SEQUENCES:
                                                                                                                                                                                                                                                      TOPOLOGY:
                                                                                                                                                                                                                                                                     STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE:
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APPLICATION NUMBER: US UE
APPLICATION 117-MAR-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/0 FILING DATE: 06-DEC-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/214,064
                                                                                                         2 LNASVTSPLLE 12
                                                                                                                                                                                                                                                                                                     LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5 VLRASVSSPFPE 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     94304
                                                                                                                                           Similarity 7; Conser
                                                                                                                                                                                                                                                                                      amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             975 California Avenue
                                                                                                                                                                                                                                                                                                 3218 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Goldbach,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Peters, Dirk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         van Grinsven J., Martinus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     E: Floppy disk
IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sandoz Agro,
                                                                                                                                                                                                                                                                     unknown
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                                                                                                                                                         61.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Robert W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US 08/032,235
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                                                                                                                                                                                                                                                                                                                                                                                                 137-1061
                                                                                                                                         Score 34; DB 1; L, Pred. No. 8.7e+02; 1; Mismatches 3;
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1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           in or Relating to Organic
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                                                                                                                                                                       Length 3218;
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NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                    nt No. 5352575.

APPLICANT: PETROVSKIS, ERIK A.; POST, LEONARD E.; TIMMINS, JAMES G.
TITLE OF INVENTION: PSEUDORABIES VIRUS PROTEIN
NUMBER OF SEQUENCES: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                FILING DATE: 20-APR-1990 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Brown, Scott A.
REGISTRATION NUMBER: 32,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION: TELEPHONE: (617) 498-8224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                                                                                                                                                                   13 IFDAAVTXPTLE 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: 1
                 APPLICATION NUMBER: 784,787 FILING DATE: 04-OCT-1985
                                                                                 APPLICATION NUMBER: 100,817 FILING DATE: 29-JUN-1987
                                                                                                                                                  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                APPLICATION NUMBER: 886,260 FILING DATE: 16-JUL-1986
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY:
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APPLICATION NUMBER:
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                                                                                                                                                                APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  amino acid
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87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 131 amino acids
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Bowman, Michael
                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Treacy, Maurice
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LaVallie, Edward
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Merberg, David
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Jacobs, Kenneth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Racie, Lisa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protein
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J. 876-5851
NO: 8:
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50.0%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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; LENGTH: 350 5352575-7

FILING DATE: 26-MAR-1986

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Query Match
Best Local Similarity
"-+-hes 8; Conserve
                                                                                                 Query Match
Best Local Similarity
6; Conserv
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RESULT 6
US-08-232-538-12
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                                                                                                                                                                                                                                                                                TELEX: 66141 PENNIE
NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   tent No.
                                                                                                                                                                                                                                                                                                                        NAME: Halluin, Albert P.
REGISTRATION NUMBER: 25,527
REFERENCE/DOCKET NUMBER: 83
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-854-3660
                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                MOLECULE TYPE:
                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
                                                        11 |:|||
400 LNGSITSP 407
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                                                                                                                                                                                                          STRANDEDNESS:
TOPOLOGY: un
                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: New York
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/0 FILING DATE: 13-DEC-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET:
                                                                                 2 LNASVTSP 9
                                                                                                                                                                                                                                                     LENGTH:
                                                                                                                                                                                                                                                                                                            TELEFAX:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION:
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                                                                                                                                                                                                                                        amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3: Pennie & Edmonds
1155 Avenue of the Americas
                                                                                                                                                                                                                                                     788 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hojima,
                                                                                                                                                                                                                                                                                                            415-854-3694
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Brenner, Mitch
                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       U.S.A.
                                                                                                                                                                                                             unknown
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Prockop, Darwin J.
                                                                                                                                                                                                                         unknown
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RECOMBINANT C-PROTEINASE AND ITS USE FOR DRUG DEVELOPMENT FOR THE TREATMENT OF DISEASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       60.0%;
72.7%;
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                                                                                                                            Score 33;
Pred. No.
                                                                                                             Mismatches
                                                                                                                                         DB 1;
                                                                                                                             5e+02;
                                                                                                                                      Length 788;
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                                                                                                             l; Indels
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ZIP: 07065-0900 COMPUTER READABLE FORM: MEDIUM TYPE: Diskette

CURRENT APPLICATION DATA:

OPERATING SYSTEM: SOFTWARE: Microsc

IBM Compatible SYSTEM: DOS Microsoft Word 6.0

COMPUTER:

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US-08-232-538-12
                                                                                                                                                                                                                              US-08-786-164-12
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                                                                                                                                                                                                                                                                                                                                                   Matches 6; Conserv
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                                                                                                                                                                                                             Sequence 12,
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                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (908) 594-4720
NEORMATION FOR SEQ ID NO: 12:
                                                                                                                                                                                                tent No.
                                                                                                     TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 661 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 18
TELECOMMUNICATION INFORMATION:
TELEPHONE: (908) 594-3905
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                    APPLICANT:
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                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                        223 VLNCTATTPL 232
                                                     ADDRESSEE: Met
                                         STREET: P.O. CITY: Rahway
                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Wallen, John WREGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: Rahway
COUNTRY: USA
ZIP: 07065-0900
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                                                                                                                                                                                                                                                                                                                     1 VLNASVTSPL 10
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                               공
                                                                                                                                                                                                             Application US/08786164
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                                                                                                                                                 KENDALL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              USA
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                                                                                                                                                                  THOMAS, KENNETH A.
                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                              linear
                                                        Merck & Co., Inc.
O. Box 2000, 126 E. Lincoln Ave
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                                                                                                                                                                                                                                                                                                                                                                   58.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RICHARD L.
INHIBITOR OF VASCULAR ENDOTHELIAL CELL
                                                                                                                      CELL GROWTH FACTOR
                                                                                                                                                     RICHARD
                                                                                                                                    INHIBITOR OF VASCULAR ENDOTHELIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        us/08/232,538
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                                                                                                                                                                                                                                                                                                                                                   2; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                 Score 32; DB 1;
Pred. No. 3.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                 Length 661;
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PRIOR APPLICATION DATA:

APPLICATION NUMBER:

CLASSIFICATION: APPLICATION NUMBER: FILING DATE: 21-JA

21-JAN-1997

US/08/786,164

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Db
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Best Local Similarity
"hhes 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 6, Application US/08232538 Patent No. 5712380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
                                                                                         TELEFAX: (908) 594-4720
NFORMATION FOR SEQ ID NO: (
                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 687 amino acids
                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE:
                                                                                                                       REFERENCE/DOCKET NUMBER: 18
TELECOMMUNICATION INFORMATION:
TELEPHONE: (908) 594-3905
                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION: NAME: Wallen, John W.III
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION: NAME: Mark Hand, J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 18
FELECOMMUNICATION INFORMATION:
TELEPHONE: 908-594-3905
 OLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       223 VINCTATTPL 232
                                                                                                                                                                          REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: sir
                          STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: USA
ZIP: 07065-0907
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REGISTRATION NUMBER: 36,545
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                                                                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Merck & Co., Inc.
STREET: P.O. Box 2000 126 E Lincoln Avenue
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                IOPOLOGY:
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              linear
protein
                            single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GROWTH FACTOR
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                                                                                                                                                                                                                                                    US/08/232,538
                                                                                                                                                                     35,403
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Pred. No. 3.2e+02;
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US-08-874-678-1 RESULT

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Sequence 1, Application US/08874678 Patent No. 5952199

GENERAL INFORMATION:

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                                                                                                                           ; MOLECULE TYPE: protein US-08-786-164-6
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Best Local Similarity
                                                        Matches
                                                                  Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION: TELEPHONE: 908-594-3905
                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: Mark Hand, J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             249 VINCTATTPL 258
249 VLNCTATTPL 258
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM Compat
OPERATING SYSTEM: DC
SOFTWARE: Microsoft
                                                                                                                                                          TOPOLOGY:
                                                                                                                                                                     STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: FILING DATE: 21-JA
                                                                                                                                                                                                                                                        TELEFAX: 908-594-4720
                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: USA
ZIP: 07065-0900
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                         1 VLNASVTSPL 10
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                                                                                                                                                                                  amino acid
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P.O. Box 2000, 126 E. Lincoln Ave
                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            THOMAS, KENNETH A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       IBM Compatible
                                                                                                                                                          linear
                                                                                                                                                                                                                                                                                                                                                                                                                 21-JAN-1997
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60.0%;
                                                                    58.2%;
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INHIBITOR OF VASCULAR ENDOTHELIAL
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                                                                    Score 32;
Pred. No.
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Pred. No.
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Query Match
Best Local Similarity
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; MOLECULE TYPE:
US-08-874-678-1
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                                                                                                                                                                                                                                                                                           quence
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NFORMATION FOR SEQ ID NO: 1:
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                                                                                                                                                                                                                                                                         tent No.
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                     CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                        APPLICANT:
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PRIOR APPLICATION DATA:
                                                                                                                                                   TITLE OF INVENTION:
                                                                                                                                                                                      APPLICANT:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 07-MAY-1996
ATTORNEY/AGENT INFORMATION:
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                                                                 STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: United ZIP: 94111-4187
                                                   CITY: San
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH:
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                                                                                   ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                           1 VLNASVTSPL 10
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5. 6100071
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Y: United States 94111-4187
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f: United States
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                               California
                                                               E: Flehr, Hohbach, Test, Albritton & Herbert Four Embarcadero Center, Suite 3400
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Four Embarcadero Center, Suite 3400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               758 amino acids
                                                                                                                                                 Ferrara, Napoleone
VENTION: NOVEL INHIBITORS OF VASCULAR ENDOTHELIAL
VENTION: GROWTH FACTOR ACTIVITY, THEIR USES AND PROCESSES FOR THEIR
                                                                                                                                                                                                      Presta, Leonard
                                                                                                                                                                                                                      Chen, Helen H.
                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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Chen, Helen H
                                                                                                                                                                                                                                      Davis-Smyth, Terri L
                                               Francisco
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60.0%;
                                                                                                                                 PRODUCTION
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Pred. No. 3.8e+02;
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US-08-643-839-1
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Best Local Similarity . 60.0%;
Matches 6; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                 TELEFAX: (908) 594-4720 INFORMATION FOR SEQ ID NO: 14:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: Wallen, John W.III
REGISTRATION NUMBER: 35,403
                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: A-TELECOMMUNICATION INFORMATION: TELEPHONE: (415), 781-1989
                                                                   TELECOMMUNICATION INFORMATION: TELEPHONE: (908) 594-3905
                                                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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LENGTH: 758 amino acid
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ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Walter H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA:
                 SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
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TELEFAX: (4.
TELEX: 910
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STRANDEDNESS: unl
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ZIP: 07065-0907
                                                                                                                                                                                                                                                                                                                                                                         CITY: Rahway
                                                                                                                                                                                                                                                                                                                                                                                           STREET:
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                                                                                                      REFERENCE/DOCKET NUMBER:
                                                                                                                                                                    CLASSIFICATION:
                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                     APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Merck & Co., Inc.
STREET: P.O. Box 2000 126 E Lincoln Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 VLNASVTSPL 10
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10 277299
780 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        unknown
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                            GROWTH FACTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Richard L. RICHARD ENDOTHELIAL CELL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Release #1.0, Version #1.30
                                                                                                                                                                                                       US/08/232,538
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Pred. No.
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TOPOLOGY:

linear

STRANDEDNESS: single

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Best Local Similarity
"~+^hes 6; Conserve
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 13
US-08-786-164-14
                                                                         Query Match
Best Local Similarity
                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 14, Appli
Patent No. 5861484
                                                                                                                                                                                                                                         NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ENERAL INFORMATION:
                                                                                                                                            STRANDEDNESS: single TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 780 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: THOMAS, KENNETH A. APPLICANT: KENDALL, RICHARD L. TITLE OF INVENTION: INHIBITOR OF VASCULAR ENDOTHELIAL TITLE OF INVENTION: CELL GROWTH FACTOR
                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
249 VLNCTATTPL 258
                                                                                                                                                                                                                                                                                   REGISTRATION NUMBER: 36,545
REFERENCE/DOCKET NUMBER: 188
CELECOMMUNICATION INFORMATION:
TELEPHONE: 908-594-3905
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: Microsoft Word 6.0 URRENT APPLICATION DATA:
                            1 VLNASVTSPL 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                UMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             249 VLNCTATTPL 258
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CODERATING SYSTEM: DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: USA
ZIP: .07065-0900
                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: FILING DATE: 21-JA
                                                                                                                                                                                                                                                                       TELEFAX: 908-594-4720
                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET:
                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 VLNASVTSPL 10
                                                           6;
                                                                                                                                                                                           : 780 amino acids amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 E: Merck & Co., Inc.
P.O. Box 2000, 126 E. Lincoln Ave.
                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                      58.2%;
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                                                                       Score 32; 1
Pred. No. 3
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Pred. No.
                                                           Mismatches
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                                                                                     Length 780;
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                                                         Indels
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RESULT 14 US-08-340-011-5

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US-08-901-710-5
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                                                                                                                                                                                                                                                                                     Sequence 5;
                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                 Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 312/474-0448
TELEX: 25-3856
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                      GENERAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                        TITLE OF
                                                                                                                                                         APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS
                                                                                           NUMBER OF SEQUENCES:
                                                                                                           TITLE OF INVENTION:
                                                                                                                                          APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: 435 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Patentin Rel
                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: 32
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312/474-6300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 09-OCT-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                256 VLNCTATTPL 265
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FITLE OF INVENTION:
             CITY: Chicago
STATE: Illinois
                                             STREET:
                                                            ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: 312/474-0448
COUNTRY: United States of America
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 07/9 FILING DATE: 09-OCT-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Meyers, Thomas C
                                                                                                                       T: Kaipainen, Arja
T: Matikainen, Marja-Terttu
INVENTION: FLT4, A RECEPTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    amino acid
                                                                                                                                                                                                                                                                                   Application US/08901710
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Application US/08340011
                                             E: Marshall; O'Toole; Gerstein, Murray & Borun 6300 Sears Tower, 233 South Wacker Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       E: Marshall, O'Toole, 6300 Sears Tower, 233
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1311 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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                                                                                                                                                                      Korhonen, Jaana
                                                                                                                                                                                                  Aprelikova, Olga
Pajusola, Katri
                                                                                                                                                                                                                                       Alitalo,
                                                                                                                                                                                        Armstrong,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     single
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                                                                                                                                                                                                                                      Kari
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FLT4, A NOVEL RECEPTOR TYROSIN KINASE 20
                                                                                                                                                                                                                                                                                                                                                                                                                                           58.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 32; DB 1;
Pred. No. 7.2e+02;
                                                                                                                        RECEPTOR TYROSINE KINASE,
                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gerstein, Murray & Borun
South Wacker Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 1311;
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                                                                                                                          AND USES
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TELEFAX: 312/*.

TELEFAX: 312/*.

TELEX: 25-3856

INFORMATION FOR SEQ ID NO: 5:

SEQUENCE CHARACTERISTICS:

LENGTH: 1311 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

US-08-901-710-5
Search completed: June 4, 2001, 12:14:28 Job time: 221 sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/901,710

FILING DATE:

CLASSIFICATION DATA:

APPLICATION NUMBER: 08/340,011

FILING DATE: 14-NOV-1994

PRIOR APPLICATION NUMBER: 08/257,754

PRIOR APPLICATION NUMBER: 08/257,754

PRIOR APPLICATION NUMBER: 09-UL-1994

PRIOR APPLICATION NUMBER: 07/959,951

FILING DATE: 09-UL-1992

APPLICATION NUMBER: 07/959,951

FILING DATE: 09-CT-1992

ATTONUMEY/AGENT INFORMATION:

NAME: GASS, DAVID A

REGISTRATION NUMBER: 38,153

REFERENCE/DOCKET NUMBER: 28113/33824

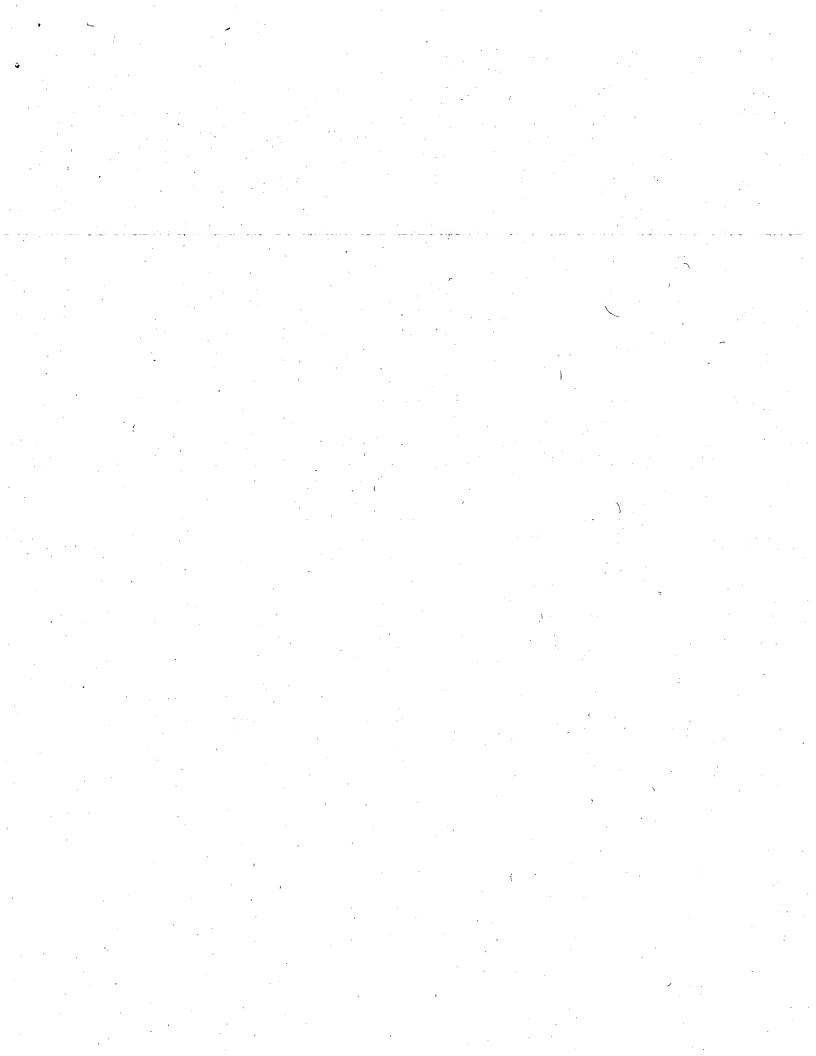
TELEPHONE: 312/474-6300

TELEFAX: 312/474-0448

TELEPAN: 312/474-0448
                                                                                                                                                                                                     Query Match 58.2%; Score 32; DB 3; 1
Best Local Similarity 60.0%; Pred. No. 7.2e+02;
Matches 6; Conservative 2; Mismatches 2
                                                                                                               1 VLNASVTSPL 10
||| : |:||
256 VLNCTATTPL 265
                                                                                                                                                                                                     2; Indels
                                                                                                                                                                                                                                               Length 1311;
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В Ş

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Compugen Ltd.

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Post-processing: Minimum Match 100%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Minimum DB seq length: 0
Maximum DB seq length: 2000000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Title:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OM protein - protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Perfect score:
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        Pred No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                            Score
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Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       June 4, 2001, 12:11:07; Search time 65.22 Seconds (without alignments) 12.645 Million cell updates/sec
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pir2:*
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hypothetical prote hypothetical prote conserved hypothet
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Fc gamma (IgG) rec
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Fc gamma (IgG) rec
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ankyrin-like prote
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alcohol dehydroger
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hypothetical prote
probable DNA-direc
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                                                                    hypothetical prote
                                                                                    tubulin-folding co
                                                                                                                         hypothetical prote
                                                                                                                                                                                                                                                                                                                                                                gamma (IgG) rec
gamma (IgG) rec
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ALIGNMENTS

A41357
Fc gamma (IgG) receptor I (high affinity)

form b -

RESULT

		1 2 2 6 1 1 1 2 2 6 1		, .
			,	÷ .
RESULT 2 A39878 RESULT 2 A39878 FC gamma (IgG) receptor I-A (high affinity) precursor - human N;Alternate names: CD64 C;Species: Homo sapiens (man) C;Date: 30-Dec:1991 #sequence_revision 06-Sep-1996 #text_change 21-Jul-2000 C;Accession: A39878; I70304; B41357; S03018; I57525 R;van de Winkel, J.G.J.; Ernst, L.K.; Anderson, C.L.; Chiu, I.M. J. Biol. Chem. 266, 13449-13455, 1991 A;Title: Gene organization of the human high affinity receptor for IgG, FcgammaRI (CD A;Reference number: A39878; MUID:91302383 A;Accession: A39878 A;Molecule type: DNA A;Residues: 1-374 <van></van>	vative 0; Mismatc 12 204	MBL:X14355; NID:g31333; PIDN:CAA32536.1; PID:g31334 MRI:X14355; NID:g31333; PIDN:CAA32536.1; PID:g31334 ranslated the codon ACT for residue 25 as Ala ma receptor I; immunoglobulin homology ve splicing; immunoglobulin receptor; transmembrane protunoglobulin homology <imm> 100.0%; Score 55; DB 2; Length 344; ty 100.0%; Pred. No. 0.0038;</imm>	A;Molecule type: mRNA A;Residues: 1-344 <all1> A;Cross-references: GB:X1435; GB:M21090; NID:g31333; PIDN:CAA32536.1; PID:g31334 A;Cross-references: GB:X1435; GB:M21090; NID:g31333; PIDN:CAA32536.1; PID:g31334 A;Clen, J.M.; Seed, B. Nucleic Acids Res. 16, 11824; 1988 Nucleic Acids Res. 16, 11824; 1988 A;Title: Nucleotide sequence of three cDNAs for the human high affinity Fc receptor (A;Reference number: S03018; MUID:89098339 A;Accession: S03019 A;Molecule type: mRNA</all1>	N;Alternate names: CD64 C;Species: Homo sapiens (man) C;Atternate names: S03019 C;Attes: 03-Apr-1992 #sequence_revision 03-Apr-1992 #text_change 21-Jan-2000 C;Accession: A41357; S03019 Science 243, 378-381, 1989 A;Title: Isolation and expression of functional high-affinity Fc receptor complementa A;Reference number: A41357; MUID:89100284 A;Accession: A41357

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submitted to the EMBL Data Library, January 1995
A;Description: Cloning and overexpression in Escherichia
A;Reference number: S51120
                                                                   A; Residues: 1-347 <CAN>
                                                                                                                               A; Accession:
                                                                                                                                                                                                                                                           R; Cannio,
                                                                                                                                                                                                                                                                                                                                                                                    alcohol dehydrogenase (EC 1.1.1.1) - Sulfolobus sp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Map position:
                                                                                                                                                                                                                                                                                                          Species: Sulfolobus sp.; Species: Sulfolobus sp.; Species: Sulfolobus sp.; Species: 19-Mar-1997 #sequence_revision 09-May-1997 #text_change 11-Jun-1999; Date: 19-Mar-1997 #sequence_revision 09-May-1997 #text_change 11-Jun-1999; Date: 19-May-1997 #text_change 11-Jun-1997 #text_change 11-Jun-1997 #text_change 11-Jun-1997 #text_change 11-Jun-1999 #text_change 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ;Molecule type: mRNA;Residues: 1-24,'S',26-337,'T',339-374 <AL2>;Coss-references: EMBL:X14356; NID:g31331; ;Perez, C.; Wietzerbin, J.; Benech, P.D.
                                                                                          Molecule type: DNA
                                                                                                                                                                                                                                                                                      Accession: S51120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Status: translated from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Accession: 170304
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cross-references: GB:M63830; GB:M63835; NID:g180277; Porges, A.J.; Redecha, P.B.; Doebele, R.; Pan, L.C.;
                              Cross-references: EMBL:Z47543;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Supertamily: Fc gamma receptor I; immunoglobulin homology Keywords: alternative splicing; glycoprotein; immunoglobu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Title: Isolation and expression of functional high-affinity Reference number: A41357; MUID:89100284 Accession: B41357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cross-references: GB:L03418; Allen, J.M.; Seed, B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Molecule type: mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Reference number:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cross-references: GDB:135911; OMIM:146760
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gene: GDB:FCGR1A; CD64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Reference number: I57525; MUID:93204964 Accession: I57525
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Molecule type: mRNA
Residues: 1-24,'S',26-337,'T',339-374 <AL1>
Cross-references: GB:X14356; GB:M21091; NID:g31331;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Residues: 1-374 <RES>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          293-313/Domain: transmembrane #status
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cross-references:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Molecule type: DNA
Residues: 1-8 <RE2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ience 243,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ccession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6-292/Domain: extracellular #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  eference number: S03018; N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    l7-170/Domain: immunoglobulin homology <IMM2>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      eic Acids Res. 16,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cession: S03018
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           78,152,159,163,195,240/Binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /Domain: signal sequence #status
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
nes 12; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 VLNASVTSPLLE 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            translated from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Biol. 13,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     378-381,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ol. 13, 2182-2192, 1993 cis-DNA elements involved in myeloid-cell-specific
                                                                                                                                                                                                                                                        Fiorentino, G.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Fc gamma receptor
source:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Seed, B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   <RE2>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1q21-1q21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           90, 2102-2109,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%;
llarity 100.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GB:S57204; NID:g298692; PIDN:AAD13887.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           I55577;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11824, 1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  e of three cDNAs
MUID:89098339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MUID: 93055454
                                                                                                                                                                                                                                                        Carpinelli, P.; Rossi, M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NID: g184840;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1992
                              NID: 9623347;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 55; DB Pred. No. 0.0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             family
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              atus predicted <TMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         gene products
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PIDN: AAA36049.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    for the human high affinity Fc receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.0042;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PIDN:CAA32537.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB
                                 PIDN:CAA87591.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         <EXT>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           <SIG>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PIDN:CAA32537.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PIDN:AAA35678.1; PID:g180279 Salmon, J.E.; Kimberly, R.P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (Asn)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             in
                                                                                                                                                                                           coli
                                                                                                                                                                                                                                                           Bartolucci,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PID:g4261587
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (covalent) #status
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PID:g292169
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PID:g31332
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                                 PID:g623348
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      receptor;
                                                                                                                                                                                           the genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         transmembrane
                                                                                                                                                                                               encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               and
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C;Function:
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                                                                   A;Cross-references: GB:AE000876; GB:AE000666; NID:g2622140;
                                                                                                                                                                                                                                                                                                                                                                                       R;Smith,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Cross-references: GB: ALO21958;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A70824
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strain

58; GB:AL123456; NID:g3261536; H37Rv

PIDN:CAA17509.1;

PID:e125

67.3%; 77.8%;

Score 37; Pred. No.

6.5;

2,

Length 175

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A; Authors: Sqares, R.; Sulston, J.E.; Taylor, K.; A; Title: Deciphering the biology of Mycobacterium A; Reference number: A70500; MUID:98295987
                                                                                                                                                                                                                                                          hypothetical glycine-rich protein RVO C; Species: Mycobacterium tuberculosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  F;38,68,154/Binding site: zinc, F;98,101,104,112/Binding site: ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       F;174-203/Region: beta-alpha-beta NAD nucleotide-binding fold F;11,213/Modified site: N6-methyllysine (Lys) #status predicted F;38,68,154/Binding site: zinc, catalytic (Cys, His, Cys) #status
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C;Keywords: alcohol metabolism; heat-stable prote;
F;23-337/Domain: long-chain alcohol dehydrogenase
A; Molecule type: DNA
A; Residues: 1-175 <COL>
                                                                                                                                                                                                 R;Cole, S.T.;
                                                                                                                                                                                                                    C; Accession: A70824
                                                                                                                                                                                                                                       C;Date: 17-Jul-1998 #sequence_revision 17-Jul-1998 #text_change 22-Oct-199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C; Superfamily: alcohol
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Pathway: alcohol
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Description:
                                 A;Status: preliminary; nucleic acid sequence
                                                        A; Accession: A70824
                                                                                                                                      Nature 393,
                                                                                                                                                       Rajandream, M.A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local (
                                                                                                                                                                               Connor, R.;
                                                                                                                                                                                                                                                                                                                                                                                   220 VINASVQDPLAE
                                                                                                                                                                                                                                                                                                                                                                                                                         VLNASVTSPLLE
                                                                                                                                      537-544
                                                                                                                                                                                                                                                                             glycine-rich protein Rv0742 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity 66.7
8; Conservative
                                                                                                                                                                               Davies,
                                                                                                                                                                                               Brosch,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    catalyzes the oxidation of primary
                                                                                                                                                     Rogers, J.; Rutter,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             degradation
                                                                                                                                                                               R.; Parkhill, J.; Garnier, T.; Churcher, C.; R.; Devlin, K.; Feltwell, T.; Gentles, S.; Ha
                                                                                                                                                                                                                                                                                                                                                                                   231
                                                                                                                                                                                                                                                                                                                                                                                                                           12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            dehydrogenase; long-chain alcohol dehydrogenase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      site: zinc,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      69.1%;
66.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       heat-stable protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 38; DB Pred. No. 9.2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        noncatalytic
                                                                                                                                                            Seeger,
                                                                                                                                                                                                                                                                                Mycobacterium tuberculosis (strain H37RV)
                                     not shown;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1;
                                                                                                   Whitehead, S.; Barrell,
tuberculosis from the c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    homology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    and secondary alcohols
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (Glu,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 347
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         homodimer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cys,
                                     translation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        <LADH>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cys,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         metalloprotein;
                                                                                                                                                                               Hamlin,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        predicted Cys) #sta
                                                                                                                                                                                                 Harris, D.;
                                          not shown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        #status predic
                                                                                                   complete
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                                                                                                                                                                                 Holroyd
                                                                                                                                                                                                      Gordon
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                                                                                                   geno
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A; Molecule type: DNA
A; Residues: 1-361 <MTH>
                                                                           A; Reference number: A; Accession: D69005
                                                                                                   A; Title: Complete genome sequence of Methanobacterium thermoautotrophicum Delta A; Reference number: A69000; MUID:98037514
                                                                                                                                                                                                                                                                                 C;Species: Methanobacterium thermoautotrophicum
C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 21-Jul-2000
                                             A; Status: preliminary; nucleic acid sequence
                                                                                                                                                                                                                                                      C; Accession:
                                                                                                                                                                                                                                                                                                                                  hypothetical protein MTH1037 - Methanobacterium thermoautotrophicum (strain Delta
                                                                                                                        ". Bacteriol. 179, 7135-7155, 1997
                                                                                                                                                                                                      Qiu, D.; Spadafora,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4 ASVTSPLLE 12
                                                                                                                                                                                                                              D.R.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity 7; Conser
                                                                                                                                                                                                                                                            D69005
                                                                                                                                                                           Doucette-Stamm, L.A.; Deloughery, C.; Lee, H.; dafora, R.; Vicaire, R.; Wang, Y.; Wierzbowski, G.M.; Daniels, C.J.; Mao, J.; Rice, P.; Noell:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ?
                                                  not shown; translation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                 Noelling,
                                                                                                                                                                                                      Dubois, J.; Aldredge, T
J.; Gibson, R.; Jiwani,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0,
                                                        not shown
                                                                                                                                                                                      Reeve,
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Ξ)

source:

PIDN: AAB85533.1;

PID: g262

Η:

fu

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A:Status: A:A-CARA A: Nolecule type: DNA
A; Molecule type: DNA
A; Mose DNA
A; Rose, D.J.; Mau, B.; Shao, Y.
Science 277, 1453-1462, 1997
A;Title: The complete genome sequence of F
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Molecule type: DNA
A;Residues: 1-1118 <NOS>
A;Cross-references: EMBL:Y11606; NID:e1033579; PID:e321611; PIDN:CAA72340.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      probable DNA-directed DNA polymerase (EC 2.7.7.7) -
C;Species: mitochondrion Pichia kluyveri
C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 밁
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C; Superfamily: hypothetical protein MTH778
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Reference number: A; Accession: T28426
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              submitted to the EMBL Data Library, March 1997 A;Reference number: Z20485
                                A;Status: nucleic
                                             A; Accession: A64809
                                                                                                                                                                                                                                                                                                                     A; Reference number:
                                                                                                                                                                                                                                                                                                                                   A; Title: Overexpression
                                                                                                                                                                                                                                                                                                                                                 Biochem. J.
                                                                                                                                                                                                                                                                                                                                                                                                                            N;Alternate
                                                                                                                                                                                                                                                                                                                                                                                                                                         transcription regulator fark,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Status: preliminary; translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             R; Nosek, J.
                                                                                                                                                                                                                             EBS Lett.
                                                                                                                                                                                                             ;Title: Identification of a
                                                                                                                                                                       Reference number: S50896; MUID:95104410
Accession: S50896
                                                                                                                                                                                                                                                                                                                                              Buck, D.; Guest, J.R. cochem. J. 260, 737-747,
                                                                                                                                                                   Status: preliminary
                                                                                                                                                                                                                                           Cross-references: EMBL:X15790; NID:g41518; Quail, M.A.; Dempsey, C.E.; Guest, J.R.
                                                                                                                                                                                                                                                                          Residues: 1-240 <B
                                                                                                                                                                                                                                                                                                                                                                                                Date:
                                                                                                                                                                                                                                                                                                                                                                                                             Species:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Accession:
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                                                                                                                                                                                                                                                                                                   Accession: S04645
                                                                                                                                                                                                                                                                                                                                                                                Accession:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Genetics
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local
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sion: S04645; S50896; A64809; PC4138
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                                                                                                                                                                                                                                                                                                                                                                                                             Escherichia coli
                                                                                                                                                                                                                                                                                                                                                                                                                            names:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          mitochondrion;
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                                                                                                                                                                                                                                                                         1-240 <BUC>
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                               acid sequence not shown; translation not shown
                                                                                                                                                                                                                                                                                                                                                                                                                         gp30
                                                                                                                                                                                                                                                                                                                      S04645;
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71.48;
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72.78;
                                                                                                                                                                                                                                                                                                                                                  1989
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                                                                                                                                                                                                                                                                                                                      site-directed MUID:89350876
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              37;
                                                                            Escherichia
                                                                                                                                                                                                                                                                                                                                   mutagenesis
                                                                                                                     C.A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB
                                                                                                                                                                                                                                                           PIDN:CAA33791.1;
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                                                                                                                     Perna,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      #text_change 15-Oct-1999
                                                                                                                                                                                                               regulator (FarR) in
                                                                                                                                                                                                                                                                                                                                                                                             #text_change 10-Sep-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ۲,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pichia kluyveri mitochondrion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 361;
                                                                            coli
                                                                                                                                                                                                                                                                                                                                   of
                                                                                                                                                                                                                                                                                                                                                                                                                                         Escherichia
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                                                                                                                       Z. T. ;
                                                                                                                                                                                                                                                                                                                                   the
                                                                                                                                                                                                                                                                                                                                   succinyl-CoA synthetase
                                                                                                                       Burland,
                                                                                                                                                                                                                                                         PID: g41519
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                         coli
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                     ٧.;
                                                                                                                                                                                                             Escherichia coli
                                                                                                                     Riley,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      linea
                                                                                                                                                                                                                                                                                                                                   of.
hypothetical protein
C; Species: Methanoco
C; Date: 13-Sep-1996
                                                                                                          밁
                                               H64489
                                                             RESULT
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220

VINASMODPLAE 231

protein MJ1521 -

Methanococcus jannaschii

revision

13-Sep-1996 #text_change 13-Sep-1998

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Biosci. Biotechnol. Biochem. 60, 3.
A;Title: Isolation and characterina.
A;Reference manufacture.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C; Function:
A; Description: probably a transcription regulator G; Superfamily: transcription regulator GntR C; Keywords: DNA binding; transcription regulation
δÃ
                                                                                                                                                                                                                                    C;Superfamily: alcohol dehydrog
C;Keywords: alcohol metabolism;
                                                                                                                                                                                                                                                                                                                  C; Complex:
C; Function:
                                                                                                                                                                                                                                                                                                                                                                     A;Cross-references: GB:S51211; NID:g261968; PIDN:AAB24546.1; PID:g261969
A;Experimental source: strain DSM1617
A;Note: sequence extracted from NCBI backbone (NCBIN:120928, NCBIP:120938)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ş
                                                                                                                                                                           F;174-203/Region: beta-alpha-beta NAD nucleotide-binding fold F;11:213/Modified site: N6-methyllysine (Lys) (partial) #status experimental
                                                                                                                                                                                                                    F;23-337/Domain: long-chain alcohol dehydrogenase homology <LADH>
                                                                                                                                                                                                                                                                         A; Description: catalyzes the oxidation of primary and secondary alcohols to aldehydes A; Pathway: alcohol degradation
                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Molecule type:
A; Residues: 1-34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Accession: A44245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Reference number: A44245; MUID:93099126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C;Species: Sulfolobus solfataricus C;Date: 10-Mar-1994 #sequence_revision
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Gene: farR; gp30
A;Map position: 16.8 min
                                                                                                                                    F;98,101,104,112/Binding site: zinc,
                                                                                                                                                                                                                                                                                                                                                         A; Note: part of this sequence was
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Title: Thermostable NAD(+)-dependent alcohol dehydrogenase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Biochemistry 31,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R; Ammendola, S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C; Accession: A44245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      alcohol
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Residues: 1-40 <UTS>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Accession: PC4138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Cross-references: GB:AE000176; GB:U00096;
                                                         Query Match
Best Local
                                                                                                                                                                                                                                    Superfamily: alcohol dehydrogenase; long-chain alcohol dehydrogenase homology Keywords: alcohol metabolism; heat-stable protein; homodimer; metalloprotein;
                                                                                                                                                                                                                                                                                                                                      Complex: homodimer
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Experimental
                                        Matches
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                                                                                                                                                             8,68,154/Binding site:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    192 LLNISQTKPILE 203
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 VLNASVTSPLLE
VLNASVTSPLLE
                                        Similarity 58. 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                  1-347 <AMM>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  source: strain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Raia, C.A.; Caruso, C.; Camardella, L.; D'Auria, S.; 12514-12523, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA
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  12
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                                                             58.3%;
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                                                                                                                                                             catalytic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               309-315,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 35;
Pred. No.
                                                           Score 35;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                         confirmed by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Kaino, Y.;
9-315, 1996
                                        red. No. 36;
Mismatches
                                                                                                                                        noncatalytic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sulfolobus solfatarious
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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                                                                                                                                                             (Cys,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NID: g1786947;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tanabe,
                                                                                                                                                                                                                                                                                                                                                         protein
                                                                               1;
                                                                                                                                                             His,
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                                                                                                                                        (Glu, Cys, Cys, Cys) #status predic
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                                          Ψ
                                                                               Length 347;
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                                                                                                                                                           Cys) #status
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H64489; White,

Blake, dek, A.

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probable phosphoglucomutase - syphilis spirochete
C;Species: Treponema pallidum subsp. pallidum (syphilis spirochete)
C;Date: 24-Jul-1998 #sequence_revision 24-Jul-1998 #text_change 05-Nov-1999
C;Accession: C71327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Authors: Kaine, B.P.; Borodovsky, M.; Klenk, H.P.; Fraser, C.P. A;Title: Complete genome sequence of the methanogenic archaeon, A;Reference number: A64300; MUID:96337999
A;Accession: H64489
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     rson, J.D.; Sadow, P.W.; Hanna, M.C.; Science 273, 1058-1073, 1996
                                                                                             κ;rraser, C.M.; Norris, S.J.; Weinstock, G.M.; White, O.; Sutton, G.G.; Dodson, rson, J.; Khalak, H.; Richardson, D.; Howell, J.K.; Chidambaram, M.; Utterback, they, L.; Weidman, J.; Smith, H.O.; Venter, J.C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 20-Apr-2000 #sequence_revision 20-Apr-2000 #text_change
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ankyrin-like protein
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A; Residues: 1-422 <BUL>
                                                                              they, L.; Weidman, J.; Smi
Science 281, 375-388, 1998
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                                                                                                                                                                                                                                                                                                                                                                         δÃ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     N; Alternate names: protein F18022
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C;Superfamily: hypothetical protein MJ0057
                          A; Reference number: A71250; A; Accession: C71327
                                       A; Title: Complete genome sequence of Treponema pallidum, the syphilis spirochete A; Reference number: A71250; MUID: 98332770
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      submitted to the Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R; Bevan, M.; Murphy, G.; Ridley, P.; Hudson, S.; Bancroft, I.; Mewes, H.W.; Rudd,
A; Status: preliminary; nucleic acid sequence not shown; translation not shown
                                                                                                                                                                                                                                                                        RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ;Bult, C.J.; White, O.; Olsen, G.J.; Zhou, L.; Fleischmann, R.D.; Sutton, G.G.; Bla
Reich, C.I.; Overbeek, R.; Kirkness, E.F.; Weinstock, K.G.; Merrick, J.M.; Glodek,
son, J.D.; Sadow, P.W.; Hanna, M.C.; Cotton, M.D.; Roberts, K.M.; Hurst, M.A.
                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 88.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Reference number: Z24493
Accession: T48596
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ;Map position:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cross-references: GB:U67593; GB:L77117; NID:g1592149;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Status: preliminary; nucleic acid sequence not shown;
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Introns: 32/3; 78/3
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Molecule type: DNA
Residues: 1-591 <BEV>
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Best Local
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7; Conserv
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58.3%;
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Pred.
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Pred. No.
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45;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           clone F18022
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McDc
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Best Local
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Local Similarity hes 6; Conserv

Conservative

63.6%;

score 35; DB; Pred. No. 91; 2; Mismatches

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Indels

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DB 1;

Length 761

1 VLNASVTSP

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S61642
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A;Residues: 1-632 <COL>
A;Cross-references: GB:AEO01219; GB:AEO00520; NID:g3322693; PIDN:AAC65401.1; PID:g332
A;Experimental source: strain Nichols
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 C;Accession: S61642; S20179; S64897; S64901; S17029
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           translation elongation factor EF-G, mitochondrial - N; Alternate names: protein L2195; protein YLR069c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Gene:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    R;Vambutas, A.; Ackerman, S.H.; Tzagoloff, A.
Eur. J. Blochem. 201, 643-652, 1991
A;Title: Mitochondrial translational-initiation and elongation factors in Saccharomyc
A;Reference number: S20177; MUID:92037620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R; Urrestarazu,
                                                            C;Superfamily: translation elongation factor G; translation elongation factor Tu homo C;Keywords: GTP binding; mitochondrion; P-loop; protein biosynthesis F;71-205/Domain: translation elongation factor Tu homology <FTU>
                                                                                                                                                                                                                     C;Genetics:
                                                                                                                                                                                                                                    A;Cross-references: EMBL:273241; NID:g1360421; PIDN:CAA97626.1; PID:g1360422; GSPDB:G
A;Experimental source: strain S288C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Molecule type: DNA
A; Residues: 1-761 <AND>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          submitted to the Protein Sequence Database, May 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Molecule type: DNA
A; Residues: 1-65, 'I',
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Accession: S2017
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R;Yambutas, A.; Ackerman, S.H.; Tzagoloff, A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Molecule type: DNA
A;Residues: 1-761 <URR>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          submitted to the EMBL Data
F;327-329/Region:
                                                                                                                   a;Map position: 12R
C;Superfamily: translation elongation factor G;
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                                                                                                                                                                                         A;Gene: SGD:MEF1; MIPS:YLR069c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Reference number: S64872
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Residues: 1-65,'I',67-232,'V',234-477,'S',479-628,'P',630-761
A;Cross-references: EMBL:X58378; NID:g3917; PIDN:CAA41267.1; PI
                                                                                                                                                                                                                                                                                                                                                          A; Reference number:
                                                                                                                                                                                                                                                                                                                                                                                                                                A; Experimental source:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Cross references: EMBL:Z73241;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Species: Saccharomyces cerevisiae
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Best Local
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                 77-84/Region: nucleotide-binding motif A (P-loop) 202-205/Region: GTP-binding NKXD motif
                                                                                                                                                                                                                                                                                                          Molecule type: DNA
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7; Conserv
                                                                                                                                                                                                                                                                                      1-761 <POH>
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GTP-binding NKXD motif GTP-binding SAK/L motif
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$64899
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   S61618
                                                                                                                                                                     SGD:S0004059; MIPS:YLR069c
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                                                                                                                                                                                                                                                                                                                                                                                                                                   41; NID:g1360421; PIDN:CAA97626.1; PID:g1360422; GSPDB:G
S288C
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Pred. No. 73;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            December 1995
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668 IMNVSVTSP 676

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Fc gamma (IgG) receptor high affinity - mouse
C; Species: Mus musculus (house mouse)
C; Date: 02-Jul-1996 #sequence_revision 02-Jul-1996 #text_change 21-Jan-2000
                                                                         RESULT
148471
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Glodek, A.; Zhou, L.; Overbeek, R.; Gocayne, J.D.; Weidman, J.F.; Nature 390, 364-370, 1997
A;Authors: Utterback, T.; Cotton, M.D.; Spriggs, T.; Artiach, P.; Smith, H.O.; Woese, C.R.; Venter, J.C.
A;Title: The complete genome sequence of the hyperthermophilic, su A;Reference number: A69250; MUID:98049343
A;Accession: C69304
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Accession: C69304
R;Klenk, H.P.; Clayton, R.A.; Tomb, J.F.; White, O.; Nelson, K.E.; Ketchum, K.A.; Dodson, Fleischmann, R.D.; Quackenbush, J.; Lee, N.H.; Sutton, G.G.; Gill, S.; Kirkness, E.F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       probable enoy1-CoA hydratase (EC 4.2.1.17) fad-1 - Archaeoglobus fulgidus
C;Species: Archaeoglobus fulgidus
C;Date: 28-May_1999 #sequence_revision 28-May-1999 #text_change 18-Jun-1999
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k; Reference number: S61996
k; Accession: S62011
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Matches 7; Conserv
                                                                                                                                                                                                                                                                                                                                                              Cross-references: GB:AE001074; GB:AE000782; NID:g2689397; PIDN:AAB90798.1; PID:g26501: Superfamily: naphthoate synthase; neny1-CoA hydratase homology Keywords: carbon-oxygen lyase; fatty acid beta-oxidation; hydro-lyase 25-175/Domain: enoy1-CoA hydratase homology <ECH>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gene: PHO85; MIPS:YPL110c Map position: 16L
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Residues: 1-1223 <SCH>
Cross-references: EMBL:U43503; NID:gl163087; PIDN:AAB68251.1;
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Residues: 1-256 <KLE>
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Whiches 8; Conserve
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                                                                                                                                                                                                                                                                                                                  A;Title: Linkage on chromosome 3 of autoimmune diabetes and defective Fc receptor for A;Reference number: I48471; MUID:93242399
A;Accession: I48471
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R; Prins, J.B.; Todd,
Science 260, 695-698,
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                                                                                                                                                                                                                                                                               A; Molecule type: mRNA
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207 VLRASVSSPFPE 218
                                   1 VLNASVTSPLLE 12
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                                                                            Conservative
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Pred. No. 55;
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Search completed: June 4, Job time: 271 sec

2001, 12:15:38

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10.944 Million cell updates/sec
93435 seqs,
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34 32 58.2 237 1 PS51 ARATH 35 32 58.2 237 1 PS52_ARATH 36 32 58.2 237 1 PS52_ARATH 37 32 58.2 237 1 PS52_SOYBN 38 32 58.2 259 1 ATP6_YEAST 39 32 58.2 259 1 ATP6_YEAST 39 32 58.2 380 1 CÓAT_CNV 39 32 58.2 428 1 FXB2_MOUSE 40 32 58.2 442 1 DZD1_XENLA 40 32 58.2 478 1 ARDE_CHLTR 41 32 58.2 478 1 ARDE_CHLTR 42 32 58.2 485 1 YC11_KLEPN 43 32 58.2 574 1 YCLF_HRSV1 44 32 58.2 574 1 GLGB_AGRTU 45 32 58.2 734 1 GLGB_AGRTU 46 32 58.2 734 1 GLGB_AGRTU 47 32 58.2 734 1 GLGB_AGRTU 48 58.2 609 1 YKD6_CAEEL 49 58.2 734 1 GLGB_AGRTU 49 58.2 734 1 GLGB_AGRTU 40 58.2 734 1 GLGB_AGRTU 41 32 58.2 734 1 GLGB_AGRTU														
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		_					xenopus	_					_	-

ALIGNMENTS

Mammalia; Eutheria; Primates; Catarrhini; Hominidae; NCBI_TaxID-9606; [1] SEQUENCE FROM N.A. MEDLINE-89098339; PubMed-2974947; Allen J.M., Seed B.; "Nucleotide sequence of three cDNAs for the human hi receptor (FcRI)."; Nucleic Acids Res. 16:11824-11824(1988). [2] [2] [2] [2] [2] [2] [2] [2] [2] [2	DARD; PRT; 374 AA. 2, Created) 2, Last sequence update) 0, Last annotation update) 0, Last annotation update) FECCIOR I (CD64). FCGI OR IGERI.) Chordata; Craniata; Vertebrata; E Primates; Catarrhini; Hominidae;
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EMBL; X14356; CAA32537.1; EMBL; X14355; CAA32536.1;

PIR; PIR; PIR; PIR;

; S03018; S03018. ; S03019; S03019. ; A41357; A41357. ; B41357; B41357.

HSSP; P126760;

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use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                  Cannio R., Fiorentino G., Carpinelli P., Rossi M., Bartolucci "Cloning and overexpression in Escherichia coli of the genes (NAD-dependent alcohol dehydrogenase from two Sulfolobus specia J. Bacteriol. 178:301-305(1996).

-i- CATALYTIC ACTIVITY: ALCOHOL + NAD(+) = ALDEHYDE OR KETONE -i- COFACTOR: BINDS FOUR ZINC IONS PER DIMER.
                                                                      the European Bioinformatics Institute.
                                                                                                              This
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                                                                                       SWISS-PROT entry is copyright. It is produced through een the Swiss Institute of Bioinformatics and the EN
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Pred. No. 0.0017;
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                                                                                                                                                                ZINC-CONTAINING ALCOHOL DEHYDROGENASE
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3C 1.1.1.1).
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Query Match
Best Local Similarity
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                                                                                                                                                                                     FEBS Lett. 416:30-34(1997).

11- FUNCTION: BINDS SPECIFICALLY AND DIRECTLY TO SUBSTRATES CONTAINING FUNCTION: BINDS SPECIFICALLY AND DIRECTLY TO SUBSTRATES CONTAINING OF IMPORT SUBSTRATES TO THE NUCLEAR PORE COMPLEX (NPC). THE COMPLEX IS SUBSEQUENTLY TRANSLOCATED THROUGH THE PORE BY AN ENERGY REQUINING, RAN-DEPENDENT MECHANISM. AT THE NUCLEOPLASMIC SIDE OF THE NPC, THE THREE COMPONENTS SEPARATE AND IMPORTIN-ALPHA AND -BETA ARE RE-EXPORTED FROM THE NUCLEUS TO THE CYTOPLASM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          035345;
15-JUL-1998 (Rel. 36, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
01-OCT-2000 (Rel. 40, Last annotation update)
IMPORTIN ALPHA-6 SUBUNIT (KARYOPHERIN ALPHA-5
                                                                                                                              -
                                                                                                                                                                                                                                                                                             alpha subunit of the nuclear pore-targeting
tissue-specific expression.";
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Eukaryota; Metazoa; (
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HSSP; P00325;
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                                                                                                                                                                                                                                                                                                        Suji L., Takumi T., Imamoto N., Yoneda Y.;
Identification of novel homologues of mouse importin alpha, the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 VLNASVTSPLLE
                                                                                                                          SUBCELLULAR LOCATION: CYTOPLASMIC.
TISSUE SPECIFICITY: ONLY SLIGHTLY DETECTED IN EHRLICH ASCITES
TUMOR CELLS, THYMUS AND SKELETAL MUSCLE, CLEARLY DETECTED IN
KIDNEY, SPLEEN, LLVER, HEART, AND LUNG. HIGH EXPRESSION IN TESTIS.
DOMAIN: CONSISTS OF AN N-TERMINAL HYDROPHILIC REGION, A
                                      SIMILARITY: BELONGS TO THE IMPORTIN SIMILARITY: CONTAINS 8 ARM REPEATS.
                                                                                        HYDROPHOBIC CENTRAL REGION COMPOSED OF 8 REPEATS, AND A SHORT HYDROPHILIC C-TERMINUS. THE N-TERMINAL HYDROPHILIC REGION CONTAINS THE IMPORTIN BETA BINDING DOMAIN (IBB DOMAIN), WHICH IS SUFFICIENT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            VINASVQDPLAE
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IE; PS00059;
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ADH_ZINC;
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Pred. No.
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Sciurognathi; Muridae;
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                                                    IMPORTIN
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AC (CATALYTIC) (BY SIMILARITY).
AC (SECOND ATOM) (BY SIMILARITY).
AC (CATALYTIC) (BY SIMILARITY).
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                                                  ALPHA FAMILY
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                                                                            NUCLEAR PROTEIN
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European Bioinformatics Institute.

There are no restrictions on ng as its content is in no

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RESULT 4
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Best Local
                                                                                   STRAIN-K12 / MG1655;

STRAIN-K12 / MG1655;

MEDLINE-97426617; PubMed-9278503;

Blattner F.R., Plunkett G. III, Bloch C.A.,

Riley M., Collado-Vides J., Glasner J.D., R.
                                                                                                                                                                                                                                                                                                                 FARR_ECOLI P13669;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               DOMAIN
SEQUENCE
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                                                                                                                                            "Overexpression and site-directed mutagenesis synthetase of Escherichia coli and nucleotide (930) that is adjacent to the suc operon."; Biochem. J. 260:737-747(1989).
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01-JAN-1990 (Rel.
01-NOV-1997 (Rel.
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Escherichia coli.
Bacteria; Proteobacteria;
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01-JAN-1990 (Rel. 13, Last seno
01-NOV-1997 (Rel. 35, Last anno
FATTY ACYL RESPONSIVE REGULATO
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                              SEQUENCE OF 1-40 FROM N.A.
                                                                             Gregor J., Davis N.W., Kirkpatrick H.A.,
                                                                                                                                                                                     Buck D.,
                                                                                                                                                                                                         SEQUENCE FROM N.A.
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PROSITE; PS50176;
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                     MEDLINE=97076625;
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                                                                                                                                                                                                                                                                                                                                                                           202
                                                                                                                                                                                                                                                                                                                                                                                             1 VLNASVTSPLL 11
                                                                   B., Shao Y.,
                                                complete genome sequence of Escherichia coli K-12."; nce 277:1453-1474(1997).
                                                                                                                                                                                                                                                                                                                                                                           VLNCSILNPLL 212
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  Horie T., Katoh A., K and characterization
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                    PubMed=9063979;
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Last annotation updat
REGULATOR (P30 PROTEI
                                                                                                                                                                                                                                                                                                                                                                                                                         . 59
                                                                                                                                                                                                                                            gamma subdivision; Enterobacteriaceae;
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Pred. No.
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527888180223C77C CRC64;
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Kaino Y., Tanabe H., Noda M.; of the heat-responsive genes
                                                                                                                                                                                                                                                                          tation update)
(P30 PROTEIN)
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                                                                              Goeden M.A.,
                                                                                      Rode C.K., Mayhew G.F.,
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                                                                                                Perna N.T.,
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                                                                                                                                                                 of the succinyl-CoA sequence of a gene
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Ol-FEB-1995 (Rel. 31, Created)
Ol-FEB-1995 (Rel. 31, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation updat
NAD-DEPENDENT ALCOHOL DEHYDROGENASE (EC 1.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       InterPro; IPR000524;
InterPro; IPR000524;
Pfam; PF00392; gntR; 1.
PRINTS; PR00035; HTHGNTR.
PROSITE; PS000043; HTH_GNTR_FAMILY; 1.
Transcription regulation; DNA-binding.
Transcription regulation; DNA-binding.
DNA_BIND 32 51 H-T-H MOTIF (BY SIMILARITY).
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Quail M.A., Dempsey C.E., Guest J.R.;
"Identification of a fatty acyl respoi
Escherichia coli.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE
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EcoGene; EG11109; farR.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        entities requires a license agreement
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEBS Lett. 356:183-187(1994)
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SIMILARITY: BELONGS TO THE GNTR FAMILY OF TRANSCRIPTIONAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FUNCTION: POSSIBLE TRANSCRIPTIONAL REGULATOR OF THE SUCCINY SYNTHETASE OPERON. BINDS SPECIFICALLY TO THE FARR PROMOTER EXHIBITS WEAK BINDING TO THE CITRATE SYNTHASE AND LIPOAMIDE
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GRHR_CLAGA STANDARD; PRT; 379 AA. 042329; 15-JUL-1998 (Rel. 36, Created) 15-JUL-1998 (Rel. 36, Last sequence update) 15-JUL-1998 (Rel. 36, Last annotation update) GONADOTROPIN-RELEASING HORMONE RECEPTOR (GNRH Clarias gariepinus (Sharptooth catfish) (Afri Eukaryota; Metazoa; Chordata; Craniata; Verte
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relationship to other alcohol dehydrogenases.";
Biochemistry 31:12514-12523(1992);
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRAIN=DSM 1617 / Aravalli R.N.;
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Ammendola S., kar.

M., Rossi M.,
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Archaea; Crenarchaeota;
NCBI_TaxID=2287;
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Intern
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STRAIN=DSM 1617 / P
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CATALTIC ACTIVITY: ALCOHOL + NADH+) - ALDEHYDE OR KETONE + NADH-
COFACTOR: BINDS FOUR ZINC IONS PER DIMER.
SUBUNIT: HOMODIMER AND HOMOTETRAMER.
SIMILARITY: BELONGS TO THE ZINC-CONTAINING ALCOHOL DEHYDROGENASE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  requires a license agreement (See http://www.isb-sib.ch/announce/an email to license@isb-sib.ch).
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    (African catfish)
Vertebrata; Eutel
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      Euteleostomi;
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RESULT 7
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Matches
YF21_METJA STANDARI
Q58916;
Q58916;
Q1-NOV-1997 (Rel. 35, C
15-DEC-1998 (Rel. 37, I
15-JUL-1999 (Rel. 38, I
PUTATIVE NA(+)/H(+) EX
                                                                                                                                                                                                             CARBOHYD
DISULFID
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                        PROSITE; PROSITE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tensen C.P., Okuzawa K., Blomenroehr M., Rebers F.E.M., Bogerd J., Schulz R.W., Goos H.J.T.; "Distinct efficacies for two endogenous ligands on a si
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Actinopterygii; Neopterygii; Teleostei; Euteleostei; Ostariophysi; Siluriformes; Clariidae; Clarias.
                                                                                                                                                                                                                                                                                                                                                                                                                                             G-protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the El
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TISSUE=Pituitary; MEDLINE=97182593;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pfam; PF00001; 7tm_1;
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                                                                                                                         1 VLNASVTSPLLE
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FUNCTION: THIS RECEPTOR MEDIATES ITS ACTION BY ASSOCIATION WITH G PROTEINS THAT ACTIVATE A PHOSPHATIDYLINOSITOL-CALCIUM SECOND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SIMILARITY: BELONGS TO FAMILY 1. OF G-PROTEIN COUPLED RECEPTORS
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PS50262; G_PROTEIN_RECEP_F1_2; 1.
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EXCHANGER MJ1521.
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58.3%;
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015131;

15-JUL-1998 (Rel.

15-JUL-1998 (Rel.

01-OCT-2000 (Rel.

IMPORTIN ALPHA-6;
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                                                                                                     HUMAN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                -!- SUBCELLULAR LOCATION: INTEGRAL MEMBRANE PROTEIN (POTENTIAL).
-!- SIMILARITY: BELONGS TO THE NA(+)/H(+) EXCHANGER FAMILY: STRONG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Science 273:1058-1073(1996).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "Complete genome sequence of the methanogenic archaeon, Methanococcus jannaschii.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Archaea; Euryarchaeota; Methanococcales; Methanococcaceae;
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Homo sapiens (Human).
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                                    (Rel.
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                                                                                      STANDARD;
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139
177
203
203
236
262
301
327
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404
45809
                        . 40, Last annotation update) SUBUNIT (KARYOPHERIN ALPHA-5 SUBUNIT).
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36, Last sequence up
40, Last annotation
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                                                                                                                                                                                                               63.6%;
58.3%;
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EFG1_YEAST RESULT

EFG1_YEAST P25039; Q99360; 01-MAY-1992 (Re

(Rel. 22, Created)

STANDARD;

PRT;

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use by non-profit institutions as 10 modified and this statement is not remo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             the expression pattern of the importin-alpha protein FEBS Lett. 417:104-108(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rochler M., Ansieau S., Prehn S., Leutz A., Haller H., Hartmann I "Cloning of two novel human importin-alpha subunits and analysis "cloning of two novel human importin-alpha protein family.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=98055463; PubMed=9395085; Ansieau S., Prehn S.,
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Mammalia; Eutheria;
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205 VLNCEILPPLLE
                           1 VLNASVTSPLLE
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DOMAIN: CONSISTS OF AN N-TERMINAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FUNCTION: BINDS SPECIFICALLY AND DIRECTLY TO SUBSTRATES CONTAINING EITHER A SIMPLE OR BIPARTITE NLS MOTIF: PROMOTES DOCKING OF IMPORT SUBSTRATES TO THE NUCLEAR PORE COMPLEX (NPC). THE COMPLEX IS SUBSEQUENTLY TRANSLOCATED THROUGH THE PORE BY AN ENERGY REQUIRING, RAN-DEPENDENT MECHANISM. AT THE NUCLEOPLASMIC SIDE OF THE NPC, THE RAN-DEPENDENT MECHANISM. AT THE NUCLEOPLASMIC SIDE OF THE NPC, THE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SIMILARITY:
SIMILARITY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DOMAIN: CONSISTS OF AN N-TERMINAL HYDROPHILIC REGION, A HYDROPHOBIC CENTRAL REGION COMPOSED OF 8 REPEATS, AND A SHORT HYDROPHILIC C-TERMINUS. THE N-TERMINAL HYDROPHILIC REGION CONTAINS THE INFORTIN BETA BINDING DOMAIN (IBB DOMAIN), WHICH IS SUFFICIENT FOR BINDING IMPORTIN BETA AND ESSENTIAL FOR NUCLEAR PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SUBUNIT: FORMS A COMPLEX WITH IMPORTIN BETA-1 SUBUNIT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              THREE COMPONENTS SEPARATE AND IMPORTIN-ALPHA AND -BETA ARE EXPORTED FROM THE NUCLEUS TO THE CYTOPLASM.
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                                                                                                                                                                                                                                                                                                                          PF01749;
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 216
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Primates;
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ASP/GLU-RICH (ACIDIC).
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InterPro; IPR000795; -.
Pfam; PF00679; EFG_C; 1.
Pfam; PF00009; GTP_EFTU;
                                                                                                                                                                                                                                                                                                                                                                                                                            EMBL; x58378; CAA41267.1;
EMBL; x94607; CAA64315.1;
EMBL; z73241; CAA97626.1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Andre B., Urrestarazu L.A.;
Submitted (MAY-1996) to the EMBL/GenBank/DDBJ databases.
-I- FUNCTION: THIS PROTEIN PROMOTES THE GTP-DEPENDENT TRANSLOCATION
OF THE NASCENT PROTEIN CHAIN FROM THE A-SITE TO THE P-SITE OF TH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Saccharomycetales; Saccharomycetaceae; Saccharomyces. NCBI_TaxID=4932;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Saccharomyces cerevisiae (Baker's yeast).
                                                                                                                                                                                                                                                                         TRANSIT
                                                                                                                                                                                                                                                                                  Transit peptide;
                                                                                                                                                                                                                                                                                               Elongation
                                                                                                                                                                                                                                                                                                                     PRINTS; PR00315; ELONGATNFCT
                                                                                                                                                                                                                                                                                                                                                                               HSSP; Pl3551; 1ELO.
SGD; S0004059; MEF1.
                                                                                                                                                                                                                                                                                                                                                                                                      PIR; S17025;
PIR; S20179;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           or send an
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the European Bioinformatics Institute.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Saccharomyces cerevisiae.";
Eur. J. Biochem. 201:643-652(1991).
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Vambutas A., Ackerman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Fungi; Ascomycota;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEF1 OR YLR069C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ELONGATION FACTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15-JUL-1998 (Rel.
15-JUL-1999 (Rel.
                                                                                                                                                                                                                                                                                               PROSITE; PS00301; EFACTOR_GTP; 1
Elongation factor; Protein biosy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                          899
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                                                                                                     Local Similarity
nes 6; Conserv
                                                                              1 VLNASVTSP 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SUBCELLULAR LOCATION: MITOCHONDRIAL SIMILARITY: BELONGS TO THE GTP-BINDI EF-G/EF-2 SUBFAMILY.
                                                          IMNVSVTSP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  and this statement is not removed. Usage by and for commercial requires a license agreement (See http://www.isb-sib.ch/announce/
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S20179.
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 STANDARD;
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38,
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84
152
205
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233
478
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                                                                                                                                                                 629
84573
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LAST ANNOTATION (MEF-G-1).
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                                                                                                                                                                 MW;
                                                                                                                                                                                      GIP (BY SIMILARITY).
GTP (BY SIMILARITY).
GTP (BY SIMILARITY).
GTP (BY SIMILARITY).
R -> I (IN REF. 1).
P -> S (IN REF. 1).
                                                                                                                Score 35; Pred. No.
                                                                                                                                                                                                                                                                        MITOCHONDRION (POTENTIAL)
  PRT;
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3E2C534509B09103
                                                                                                     Mismatches
 367 AA
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Matches
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus NCBI_TaxID-10090; [1]
                                                                                                         MOUSE STANDARD; PRT; 404 AA.
FCG1_MOUSE STANDARD; PRT; 404 AA.
P26151;
01-MAY-1992 (Rel. 22, Created)
01-MAY-1992 (Rel. 22, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
15-JUL-1999 (Rel. 38, Last annotation update)
16-MAY-1992 (Rel. 38, Last annotation update)
17-MULLING ANNAL FC RECEPTOR I PRECURSOR (FC-GAMMA)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        066809;
30-MAY-2000 (Rel. 39,
30-MAY-2000 (Rel. 39,
30-MAY-2000 (Rel. 39,
                                                                                                                                                                                                                                                                                                                                                                                                                                             PRINTS; PRO0423; CELLDVISETSZ.

PROSITE; PS01134; FTSZ_1; 1.

PROSITE; PS01135; FTSZ_2; 1.

Cell division; Septation; GTP-binding.

Cell division; Septation; GTP-binding.

OP_BIND 100 GTP (POTE)

OPT (POTE)
                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE
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modified and this statement is not removed:
entitles requires a license agreement (See l
or send an email to license@isb-sib.ch)
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Bacteria; Aquificales;
                                                             Mus musculus (Mouse).
                                                                                FCGR1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              the European Bioinformatics Institute.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Deckert G., Warren P.V., Gaasterland T., Young W.G., Lenox A.L., Graham D.E., Overbeek R., Snead M.A., Keller M., Aujay M., Huber Feldman R.A., Short J.M., Olson G.J., Swanson R.V.; "The complete genome of the hyperthermophilic bacterium Aquifex
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FTSZ OR AQ_525.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nature 392:353-358(1998).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE=98196666; PubMed=9537320;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pfam; PF00091; tubulin;
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                                                                                (FCRI) (IGG FC RECEPTOR I).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BEGIN BINDS TO AND HYDROLYZES GTP (BY SIMILARITY), SUBUNIT: AGGREGATE TO FORM A RING-LIKE STRUCTURE (BY SIMILARITY), SUBCELLULAR LOCATION: CYTOPLASMIC. ASSEMBLE AT THE INNER SURFACE OF THE CYTOPLASMIC MEMBRANE (BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FUNCTION: THIS PROTEIN IS ESSENTIAL TO THE CELL-DIVISION PROCESS. ITS SEEMS TO ASSEMBLE INTO A DYNAMIC RING ON THE INNER SUFFACE OF THE CYTOPLASMIC MEMBRANE AT THE PLACE WHERE DIVISION WILL OCCUR, AND THE FORMATION OF THE RING IS THE SIGNAL FOR SEPTATION TO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SWISS-PROT entry is copyright. It is produced through een the Swiss Institute of Bioinformatics and the EM
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87.5%;
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Pred. No.
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27;
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Best Local Similarity
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P45815;
01-NOV-1995
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CARBOHYD
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CARBOHYD
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-!- SUBCELLULAR LOCATION: TYPE I MEMBRANE PROTEIN.
-!- TISSUE SPECIFICITY: MACROPHAGE-SPECICIC.
-!- SIMILARITY: CONTAINS 3 IMMUNOGLOBULIN-LIKE C2-TYPE DOMAINS.
-!- CAUTION: IT IS UNCERTAIN WHETHER MET-1 OR MET-10 IS THE INITIATOR.
                                                                                                   _YARLI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        between the Swiss Institute of Bioinformatics and the the European Bioinformatics Institute. There are no rest use by non-profit institutions as long as its content
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                                       01-NOV-1995
01-OCT-1996
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Yarrowia lipolytica (Candida lipolytica)
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A46480; A46480.
; P12319; 1ALT.
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                                                                                                                                                                                               Conservative
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                                                                                       STANDARD;
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Tate B., McKenzie I.F.C.,
                           PROTEIN CRF1
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                                                                                                                                                                                                 1;
                                                                                                                                                                                                            Score 34;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                        CYTOPLASMIC (POTENTIAL).
IG-LIKE C2-TYPE DOMAIN (TRUNCATED).
                                                                                                                                                                                                                                                                                                                                 IG-LIKE C2-TYPE DOMAIN
IG-LIKE C2-TYPE DOMAIN
IG-LIKE C2-TYPE DOMAIN
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POTENTIAL.
                                                                                                                                                                                                                                                                                                                                                                                                            HIGH AFFINITY IMMUNOGLOBULIN RECEPTOR I.
                                                                                      PRT;
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                                                                                                                                                                                                  Mismatches
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of the mouse high affinity Fc
                                                                                      411
                                      update)
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Query Match
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GLE1_YEAST
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Pfam; PF00649; Copper Pfam; PF00649; COPPERFIST.

PRINTS; PR00617; COPPERFIST.1; 1.

PROSITE; PS00119; COPPERFIST.2; 1.

PROSITE; PS50073; COPPERFIST.2; 1.

COPPERFIST.2; 1.

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PROSITE; PS50073; COPPERFIST.2; 1.
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SEQUENCE
                                                                  STRAIN-S288C / FY1679;
Bahr A., Moeller-Rieker S., Hankeln T., Kraemer C., Schmidt E Bahr A., Moeller-Rieker S., Hankeln T., Kraemer C., Schmidt E Submitted (JUL-1996) to the EMBL/GenBank/DDBJ databases.
-I- FUNCTION: RNA EXPORT FACTOR. THE MECHANISM MAY INVOLVE TH TARGETING OF RNA/RNP MOLECULES TO OR THROUGH THE NUCLEAR COMPLEX BY MEANS OF RIPI AND NUP100 INTERACTIONS.
-I- SUBCELLULAR LOCATION: NUCLEAR. LOCALIZED PREDOMINANTLY AT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                use by non-profit institutions as modified and this statement is not rementities requires a license agreement
This SWISS-PROT entry is copyright.
                                                                                                                                                                                                                                            Murphy R., Wente S.R.; "An RNA-export mediator with an essential nuclear export signal."; Nature 383:357-360(1996).
                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A., MEDLINE=96404361; P
                                                                                                                                                                                                                                                                                                                                                                                                Saccharomyces cerevisiae (Baker's yeast).
Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
                                                                                                                                                                                                                                                                                                                                                                                                                                      GLE1 OR BRR3 OR YDL207W OR D1049
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between the Swiss Institute of Bioinformatics
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    -!- SUBCELLULAR LOCATION: NUCLEAR (POTENTIAL).
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cetales; Dipodascaceae; Yarrowia.
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POLY-GLN.
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Best Local
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P87035;
NP_BIND
                                                                                                                                                                                                                     "G proteins in Ustilago maydis: transmission of multiple signals?
EMBO J 16:1934-1942(1997).

-i- PUNCTION: GDARNINE NUCLEOTIDE-BINDING PROTEINS (G PROTEINS) AR
INVOLVED AS MODULATORS OR TRANSDUCERS IN VARIOUS TRANSMEMBRAN
SIGNALING SYSTEMS.

-i- SUBUNIT: G PROTEINS ARE COMPOSED OF 3 UNITS (ALPHA, BETA & GA
THE ALPHA CHAIN CONTAINS THE GUANINE NUCLEOTIDE BINDING SITE.
                                                                                                                                                                                                                                                                                                                                                                                                               ustilago maydis (Smut fungus).
Eukaryota; Fungi; Basidiomycota; Ustilaginomycetes;
Ustilaginomycetidae; Ustilaginales; Ustilaninaconomycetes;
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                       Pfam; PF00503; G-alpha; 1.
PRINTS; PR00318; GPROTEINA.
GTP-binding; Transducer; M
                                                                            EMBL; U85778; AAC49727.1; HSSP; P10824; 1AS3.
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01-OCT-2000 (Rel. 40, Last annotation update)
GUANINE NUCLEOTIDE-BINDING PROTEIN ALPHA-4 SU
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01-NOV-1997 (Rel.
01-OCT-2000 (Rel.
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Multigene family.
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GTP (BY SIMII
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L->A: PARTIAL LOSS OF ACTIVITY.
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L->A: PARTIAL LOSS OF ACTIVITY.
L->A: PARTIAL LOSS OF ACTIVITY.
L->A: TEMPERATURE SENSITIVE.
L->A: TEMPERATURE SENSITIVE.
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Q10197; P78749;
01-OCT-1996 (Rel. 34, Crea
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15-JUL-1999 (Rel. 38, Last
TUBULIN-FOLDING COFACTOR D
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TRANSMEM
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                                                                                                                                                                                                                                                                                                                             Wood V., Rajandream M.A., Barrell B.G., Beck A., Reinhardt R.;
Submitted (SEP-1998) to the EMBL/GenBank/DDBJ databases.
-i- FUNCTION: HAS A FUNCTION IN THE FOLDING OF BETA-TUBULIN; MAY PLAY
A VITAL ROLE IN MICROTUBULE-DEPENDENT PROCESSES AS A MICROTUBULE-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SCHPO
                     CARBOHYD
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Wood V., Ra
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MEDLINE=98119809; P
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Eukaryota; Fungi; Ascomycota; Schizosaccharomycetes;
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"Essential role of tubulin-folding cofactor D in microtubule assembly and its association with microtubules in fission yeast.";
EMBO J. 17:658-666(1998).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Schizosaccharomyces.
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P87883 hepatitis c
Q9qbc3 yaba monkey
Q9nq09 homo sapien
Q91ya6 arabidopsis
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p79119 bos taurus
Q99645 homo sapien
O53805 mycobacteri
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C21376 pichia kluy
Q91tr5 arabidopsis
Q9nmm0 leishmania
Q9nm18 leishmania
Q9nm18 leishmania
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 Q92495
                                                                                                                               SEQUENCE
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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PFAM; PF00047; ig; 3.
PRODOM; PD002534; -; 1;
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J. Clin. Invest. 90:2102-2109(1992).
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Porges A.J., Redecha
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Pred. No. 0.0094;
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                        STRAIN-IZUMO; Makashima N., Noda H.; Makashima N., Noda H.; "Nucleotide and deduced amino aci reovirus; a putative member of th Submitted (MAR-1995) to the EMBL/
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Viruses; dsRNA viruses; Reov
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                                     sequences of Nilaparvata lugens
genus Fijivirus.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eichbaum
                                                                                                   unclassified Reoviridae
                                                                                                                                                                  1199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eichbaum
                                                                                                                                                                                                                                                                                                                                                                                                                                                          se elements in a novel human Fc characterization of the gene
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                             databases
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                                                                                                                                                                                                                                                                 Length 375;
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                                                                                                                                                                                                                                                                                                                                                 "Characterization.";
rich repeat core protein.";
rich repeat core protein.";
- piol Chem. 272:18709-18717(1997).
- Piol Chem. 272:18709 A ROLE IN BONE FORMATION, AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-MAY-1997 (TrEMBLrel. 03, 01-MAY-1997 (TrEMBLrel. 03, 01-OCT-2000 (TrEMBLrel. 15, SMALL CHONDROITIN/DERMATAN S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              P79119;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           putative member of the genus Fi
J. Gen. Virol. 77:139-146(1996)
EMBL; D49694; BAA08543.1;
                                                                                                                                                              PFAM; PF00560; LRR; 3. PFAM; PF01462; LRRNT;
                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=97373567; PubMed=9228042;
                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A., PARTIAL SEQUENCE, TISSUE=FETAL EPIPHYSEAL CARTILAGE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bos taurus (Bovine)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nakashima N., Koizumi M., Watanabe H., Hiroaki N., "Complete nucleotide sequence of the Nilaparvata l putative member of the genus Fijivirus.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=96145144; PubMed=8558122;
                                                                                                                             SIGNAL
                                                                                                                                                   Proteoglycan;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=9913;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bovidae; Bovinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa;
                                                                                                                                          Connective
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                                                                                                                                                                                                                                                                                                                                                                                                           Johnson H.J.,
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                                                                                                                                                                                    NTERPRO;
                                                                                                                                                                                             INTERPRO; IPR000372; -
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                                                                                                                                                                                                                     (POTENTIAL).
SIMILARITY: THE REPEATED LEUCINE-RICH MANY PROTEINS. NUMBER IN THIS PROTEIN:
                                                                                                                                                                                                                                                              PTM: A LONG AND A SHORT FORM; PRESENT IN APPROXIMATELY EQUIMOLAR AMOUNTS; MAY ARISE BY PROTEOLYSIS OR CLEAVAGE BY EXOPEPTIDASES. PTM: THE THREE GLYCOSAMINOGLYCAN CHAINS ARE DERMATAN SULFATE
                                                                                                                                                                                                                                                                                              FLATTENED CHONDROCYTES OF THE DEVELOPING DEVELOPMENTAL STAGE: EMBRYO.
                                                                                                                                                                                                                                                                                                                     SUBCELLULAR LOCATION: EXTRACELLULAR MATRIX (BY SIMILARITY). TISSUE SPECIFICITY: PREFERENTIALLY EXPRESSED IN THE ZONE OF
                                                                                                                                                                                                                                                                                                                                           ORGANIZATION.
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Similarity 66.7%;
8; Conservative
                                                                                                                                                                                    IPR001611; -.
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31
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, Last annotation update)
SULFATE PROTEOGLYCAN PRECURSOR (PG-LB)
                                                                                                                                                   Signal; Repeat;
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                                                                                 PROTEOGLYCAN.
7 LEUCINE-RICH TANDEM REPEATS.
                                                                                                       SMALL CHONDROITIN/DERMATAN SULFATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRT;
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(INCOMPLETE).
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27;
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Best Local
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Q99645;
01-MAY-1997;
01-MAY-1997;
01-OCT-2000;
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MIM; 601657
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-I- FUNCTION: MAX HAVE A ROLE IN BONE FORMATION, AND ALSO IN
-ESTABLISHING THE ORDERED STRUCTURE OF CARTILAGE THROUGH MATRIX
ORGANIZATION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A., AND TISSUE SPECIFICITY.
TISSUE-CHONDROCYTES;
MEDLINE-97131519; PubMed-8975717;
Deere M., Johnson J., Garza S., Harrison W.R., Yoon S.-J.,
Elder F.F.B., Kucherlapati R., Hook M., Hecht J.T.;
"Characterization of human DSPG3, a small dermatan sulfate
                                                                                                                                                                                                                         SIGNAL
                                                                                                                                                                                                                                         Proteoglycan;
Connective tis
                                                                                                                                                                                                                                                                                     PFAM; PF00560; LRR; 3. PFAM; PF01462; LRRNT; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-MAY-1997 (TYEMBLIEL 03, Created)
01-MAY-1997 (TYEMBLIEL 03, Last seq
01-CCT-2000 (TYEMBLIEL 15, Last ann
SMALL CHONDROITIN/DERWATAN SULFATE P
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DSPG3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NCBI_TaxID=9606
                                                                                                                                                                                                                                                                                                                                   (NTERPRO;
                                                                                                                                                                                                                                                                                                                                                        INTERPRO;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13 ILDAAVTAPTLE 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                       MANY PROTEINS. NUMBER IN THIS PROTEIN: 6. SIMILARITY: TO OTHER SMALL CHONDROITIN/DERMATAN SULFATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                            SIMILARITY: THE REPEATED LEUCINE-RICH (LRR) SEGMENT IS FOUND MANY PROTEINS. NUMBER IN THIS PROTEIN: 6.
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                                                                                                                                                                                                                                                                                                                                                                                                                           PROTEOGLYCANS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (POTENTIAL).
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                                                                                                                                                                                                                                                                                                                                 IPR000372; -. IPR001611; -.
                                                                                                                                                                                                                                       tissue;
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                                                                                                                                                                                                                                                                Glycoprotein;
                                                                                                                                                                                                                                                                                                                                                                                                    AAC50945.1;
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Lrel. 15, Last annotation update)
DERMATAN SULFATE PROTEOGLYCAN PRECURSOR (PG-LB)
(DERMATAN SULFATE PROTEOGLYCAN 3) (DSPG3).
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58.3%;
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                                                                                                                                                 SMALL CHONDROITIN/DERMATAN SULFATE PROTEOGLYCAN.
                                                                                                                                                                                                POTENTIAL
                                                                                                                                                                                                                  POTENTIAL.
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                                                                                                                                                                                                                                                              ignal; Repeat;
                                                                                                                            LEUCINE-RICH TANDEM REPEATS
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No. 18;
                                                                                                        (INCOMPLETE).
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                                                                                                                                                                                                                                                                Leucine-repeat;
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CARBOHYD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cole S.T., Brosch R., Parkhill J., Garnier T., Churcher C., Harris Gordon S.V., Eiglmeier K., Gas S., Barry C.E. III, Tekaia F., Badcock K., Basham D., Brown D., Chillingworth T., Connor R., Davies R., Devlin K., Feltwell T., Gentles S., Hamlin N., Holroyd Hornsby T., Jagels K., Krogh A., McLean J., Moule S., Murphy L., Oliver S., Osborne J., Quall M.A., Rajandream M.A., Rogers J., Rutter S. Seener K. Skelton S. Schares S. Schares R. Steleton J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    053805;
                                                                                                                                                                                                                                                                                                                                                                                 INTERPRO; IPR000084;
PFAM; PF00934; PE; 1
PRODOM; PD001223; -;
                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 complete genome sequence.*;
Nature 393:537-544(1998).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rutter S., Seeger K., Skelton S., Squares S., Sqares R., Sul
Taylor K., Whitehead S., Barrell B.G.;
"Deciphering the biology of Mycobacterium tuberculosis from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRAIN-H37RV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bacteria; Firmicutes; Actinobacteria; Actinobacteridae; Actinomycetales; Corynebacterineae; Mycobacteriaceae; Mycobacterium.NCBI_TaxID=1773;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=98295987; PubMed=9634230;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
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                                                                                                                                                                                96 ASITSPLLD
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E42E7FC4F27C858A CRC64;
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BY SIMILARITY.
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Q9LTR5
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J. Bacteriol. 179:7135-7155(1997).
EMBL; AE000876; AAB85533.1;
INTERPRO; IPRO00130;
IPROSITE: PS00142; ZINC_PROTEASE; UNKNOWN_1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Aldredge T., Bashirzadeh R., Blakely D., Cook R., Gilbert K., Harrison D., Hoang L., Keagle P., Lumm W., Pothier B., Qiu D., Spadafora R., Vicare R., Wang Y., Wierzbowski J., Gibson R., Jiwani N., Caruso A., Bush D., Safer H., Patwell D., Prabhakar McDougall S., Shimer G., Goyal A., Pietrovski S., Church G.M., Daniels C.J., Mao J.-I., Rice P., Nolling J., Reeve J.N.;
                                                                                                                                                                                                                                                                                                                                                                                                                                   Blaisonneau J., Nosek J., Fukuhara H.; Submitted (MAR-1997) to the EMBL/GenBank/DDBJ EMBL, Y11606; CAA72340.1; -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-JAN-1998 (TrEMBLrel. 01-MAY-2000 (TrEMBLrel.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRAIN-DELTA H;
MEDLINE=98037514; PubMed=9371463;
                                                                                                                                                                                                                                                                                                                                                                                   INTERPRO; IPRO02064; -.
PROSITE; PS00116; DNA_POLYMERASE_B; UNKNOWN_1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRAIN=CBS 7907;
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01-JAN-1998 (TrEMBLrel.
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Pred. No.
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Pred.
.PRT;
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Q9NML8;
Q9NML8;
01-OCT-2000
01-OCT-2000
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01-OCT-2000 (TREMBLIEL 15,
01-OCT-2000 (TREMBLIEL 15,
PROBABLE ADA2-LIKE PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          features of the regions of (clones,");
DNA Res. 7:131-135(2000).
EMBL; AB024036; BAB02808.1;
SEQUENCE 1424 AA; 154737
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01-0CT-2000 (TrEMBLrel. 15, Last sequence update)
01-0CT-2000 (TrEMBLrel. 15, Last annotation update)
SIMILARITY TO BACTERICIDAL PERMEABILITY-INCREASING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Q9LTR5;
01-OCT-2000
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Eukaryota; Euglenozoa;
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                                                                                                                                                                         SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRAIN=COLUMBIA;
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                                                                                                                                                                                                                                                        NCBI_TaxID=5664;
                                                                                                                                                                                                                                                                                      LM26.72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "Structural analysis of Arabidopsis thaliana chromosome 3. features of the regions of 4,504,864 bp covered by sixty P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=3702;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nakamura
                                                                                                                                                                                                                                                                                                                                                                                                    483 VLNMSVTSP 491
                                                                                                                                                                                                                                     EQUENCE FROM N.A.
                                                                                 29
                                                                                                                                                                                                                                                                                                                                                                    .10
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                                                                                                   1 VLNASVTSPLLE
                                                                                 VLGANVTHPLMQ
                                                                                                                                                                                                                                                                                                                                                                                                                                         8; Conserv
                                                                                                                      Similarity 58.37; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nakamura
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Wakamura Y.,
(FEB-1999)
                                                                                                                                                                         52
                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                              PRELIMINARY;
                                                                                                                                                                         AA;
                                                                                 40
                                                                                                   12
                                                                                                                                                                         5639 MW;
                                                                                                                                 65.5%;
58.3%;
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88.9%;
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to the E
                                                                                                                                                                                                                                                                  Kinetoplastida;
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                                                                                                                                                                                                                                                                                                (FRAGMENT)
                                                                                                                                                                                                     D., Rajandream M., EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                         Last sequence update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MW;
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••
                                                                                                                                                                                                                                                                                                                              Created)
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                                                                                                                       Score 36; DB Pred. No. 7; 3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 37; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       T.,
                                                                                                                                                                         OBD2006CA2F21EC5 CRC64
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                                                                                                                                                                                                                                                                                                                                                  52
                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 10;
1.3e+02;
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                                                                                                                                           DВ
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                                                                                                                                                                                                      Ivens A., databases.
                                                                                                                                          Length 52
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                                                                                                                        Indels
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e; eurosids II;
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                                                                                                                                                                                                                 Barrell
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(TrEMBLrel. 15, (TrEMBLrel. 15,

Created)
Last sequence update)

01-OCT-2000 (TremBLrel. 15, Last POSSIBLE MICROTUBULE-ASSOCIATED |

PROTEIN (FRAGMENT).

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RESOLUT

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OROSOLu
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Best Local Similarity
---has 7; Conserv
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Best Local S
Matches 7
                                                                                           01-OCT-2000 (TremBLrel. 15, C. 01-OCT-2000 (TremBLrel. 15, L. 01-OCT-2000 (TremBLrel. 15, L. DJ622L5.1 (IMPORTIN ALPHA 7 (K. DJ622L5.1.
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01-AUG-1998
01-JUN-2000
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Murphy L., Quail M., Harris
Submitted (JUL-2000) to the
EMBL; AL160493; CAB97692.1;
NON_TER 1 1
                                                                                                                                                                               Q9NU85;
01-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Mammalia; Eutheria;
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
NCBI_TaxID-9606;
[1]
                                                                                                                                                                                                                         Q9NU85
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PFAM; PF01749;
SEQUENCE 536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Christiansen M., Hart
Submitted (APR-1998)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Leishmania
Eukaryota;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INTERPRO; IPRO02652; -. PFAM; PF00514; Armadillo_seg;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                060684;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INTERPRO;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MPORTIN ALPHA 7
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                                                                                                                                                                                                                                                                                                                         205 VLNCSILNPLL 215
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                                                                                                                                                                                                                                                                                                                                                                1 VLNASVTSPLL 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 tted (APR-1998) to the EMBL/GenBank/DDBJ databases. AF060543; AAC15233.1; -. Q02821; 1BK5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity 58.3
7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                major.
Euglenozoa;
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(TremBLrel. 07, Last sequence update)
(TremBLrel. 14, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRELIMINARY;
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AA;
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Primates;
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63.6%;
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                                                                                                                    Created)
Last sequence update)
Last annotation update)
(KARYOPHERIN)).
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EMBL/GenBank/DDBJ databases.
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                                                                                                                                                                                                                                                                                                                                                                                                                         Score 36;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 36;
Pred. No.
                                       Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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18;
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Best Local S
Matches 7
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Best Local
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01-MAY-1997
01-MAY-2000
                                                                                                                    01-MAY-1997
01-MAY-1997
01-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Submitted (FEB-2000) to the EMBL/GenBank/DDBJ EMBL; AL049795; CAB75614.1; -. SEQUENCE 566 AA; 63004 MW; 48657248DFF2678
                                                                                                                                                                                                                                                                                                                                                    epitope of hepatitis C virus in two haemophilic patients."; J. Gen. Virol. 78:583-590(1997).
EMBL; Z84308; CAB06360.1; -.~
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hepatitis C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Frankland
                 STRAIN=PATIENT 1, 1994;
MEDLINE=97201623; PubMed=9049409;
                                       SEQUENCE FROM N.A.
                                                                    Viruses; ssRNA positive-strand Hepacivirus
                                                                                                                                                                                                                                                                                                                                                                                             STRAIN-PATIENT 2, 1989;
MEDLINE-97201623; PubMed-9049409;
Devereux H.L., Brown D., Dusheiko
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Viruses; ssRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  P87859
           Devereux H.L.,
                                                                                                            PARTIAL
                                                                                                                                                   P87883;
                                                                                                                                                             P87883
                                                                                                                                                                                                                                                                                                      SEQUENCE
                                                                                                                                                                                                                                                                                                                          NON_TER
                                                                                                                                                                                                                                                                                                                                   PFAM; PF01006; HCV_NS4a; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hepacivirus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         P87859;
                                                           NCBI_TaxID=11103;
                                                                                         Hepatitis C
                                                                                                                                                                                                                                                                                                                 NON_TER
                                                                                                                                                                                                                                                                                                                                              INTERPRO; IPR000745;
                                                                                                                                                                                                                                                                                                                                                                                      "Long-term evolution
                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=11103;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    235 VLNCSILNPLL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 VLNASVTSPLL
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                                                                                                                                                                                                                                 LNASVTSPL 10
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                                                                                                             NS4
                                                                                                                                                                                                                                                     Similarity 7; Conserv
                                                                                                         7 (TrEMBLIEL. 03, C
7 (TrEMBLIEL. 03, I
1 (TrEMBLIEL. 13, I
1 GENE (PATIENT 1,
                                                                                                                                                                                                                                                                                                      44 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (Trembirel. 03, (Trembirel. 03, I) (Trembirel. 13, I) (Trembirel. 13, I) GENE (PATIENT 2,
                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                             PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRELIMINARY;
                                                                                                                                                                                                                                                                                                                 44
                                                                                                                                                                                                               12
           Brown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   positive-strand viruses,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    245
                                                                                                                                                                                                                                                                                                      4297
 D., Dusheiko of the 5'UTR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   63.6%;
                                                                                                                                                                                                                                                             63.68;
                                                                                                                                                                                                                                                                                                                                                                                     the 5'UTR and a
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Last sequence update)
Last annotation update)
, 1989, CLONE 4) (FRAGMENT).
                                                                                                          Last sequence update)
Last annotation updat
, 1994, CLONE 3) (FRAG
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                                                                                                                                        Created)
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                                                                                                                                                                                                                                                             Score 35; I
Pred. No. 9
                                                                                                                                                                                                                                                                                                      7B755D858C4A100E CRC64;
                                                                                                                                                              PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRT;
                                                                              viruses,
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G.M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   36;
No.
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79;
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9.3
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Emery V.C., La
region of NS4
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                                                                              DNA
                                                                                                                                                                                                                                                                         14;
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                                                                                                            update)
(FRAGMENT)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ņ
                                                                              stage;
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                                                                                                                                                                                                                                                                        Length 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
           Lee
                                                                                                                                                                                                                                                                                                                                                                                              Lee C.A.;
                                                                                                                                                                                                                                                                                                                                                                                      containing
                                                                                Flaviviridae;
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Result
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Maximum DB
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Perfect score:
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Maximum Match 100%
Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Scoring table:
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                                                                                                                                                                                                                                                      Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  seq
      55555555555
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length:
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61
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Gapop 10.0 , Gapext.0.5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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/SIDS6/9cgdata/geneseq/geneseqp/AA1994.DAT:
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                                                                                                                                    W60554
Y33183
                                                                                          796183
                                                                                                                        R22549
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/geneseqp/AA1992.DAT:
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                                                                                                                                                                             Description
                                                                                                                                                  Oligopeptide from
                                              Human
                                                             Human
                                                                           Human macrophage-s
                                                                                       Human macrophage-s
                                                                                                        Human
                                                                                                                    Human
                              FCRI (CDNA C
                                            macrophage-s
FCRI (CDNA c
                                                                                                        FCRI (CDNA C
                                                                                                                    macrophage-s
                                                                                                                                     sFc-gammaRI
receptor
                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT
W60554
ID W6
XX
AC W6
XX
                                Identifying peptide(s) binding specifically to protein target - by expressing on phage surface and testing for binding to immobilised oligo:peptide derived from the target, useful for, e.g. identifying specific antibodies
                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                  W60554;
                                                                        WPI; 1998-240964/21.
                                                                                                                                                                                             07-OCT-1997;
                                                                                                                                                                                                                                                      WO9815833-A1
                                                                                                                                                                                                                                                                                                                                                                                                     18-AUG-1998
                                                                                                    De Kruif CA, Logtenberg
                                                                                                                                  (UYUT-) RIJKSUNIV UTRECHT.
                                                                                                                                                                08-OCT-1996;
                                                                                                                                                                                                                          16-APR-1998
                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                96EP-0202791
                                                                                                                                                                                             97WO-NL00557
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ALIGNMENTS

Extracellular domain; CD64; identification; antibody; immunohistochemical; immunofluorescent analysis; detecell transformation; mutation; anti; oncogene Oligopeptide from extracellular domain of CD64 W60554 standard; peptide; 12 AA analysis; detection;

surface of a replicable display package, synthesising oligopeptides derived from the target protein on a solid phase, and testing for binding between the peptide and oligopeptides. The method is used to screen large peptide libraries, especially to detect antibodies, or their fragments, that bind to cell markers or that can differentiate between different that be compared to the control of the Synthetic oligopeptides W60537-61 are derived from the extracellular domain of CD64. They were synthesised on 25 polyethylene rods as adjacent, non-overlapping sets. Peptide specific phages were isolated affinity binding to the CD64-covered rods. The specification describe forms of the same protein, including bispecific antibodies that bind to two non-overlapping epitopes on the same monomeric antigen or two epitopes on different molecules. The genes/oligonucleotides that encode selected peptides can be isolated and used for recombinant production of antibodies. Selected antibodies are used, e.g. for immunohistochemical or method for the identification of a peptide able to bind specifically to target protein. The method comprises displaying the peptide on the Example 1; Page 29; 40pp; English. Sequence antibodies. Selected antibodies are used, e.g. immunofluorescent analysis, and also to detect mutation 12 A, (anti)oncogenes The specification describes cell transformation caused isolated by

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밁 Query Match Best Local S Matches 12 Human sFc-gammaRI protein fragment Y33183 standard; N l Similarity 12; Conser 100.0%; llarity 100.0%; Conservative ((first entry) Protein; 261 0; Score 61; DB 19; Pred. No. 0.00032; AA. Mismatches Indels Length 12; 0; Gaps

anti-inflammatory; drug design; therapy; inhibitor; atomic tissue damage; Fc receptor; Fc-gammaRIIa; human; FCR; inflammatory coordinate; modulator; Fc-gammaRIa; hypersensitivity; bioactive compound design; computer-assisted; py; inhibitor; Fc-gammaR; Fc-epsilonR; IgG; Ig rc-gammaR; Fc-epsilonR; IgG; IgE; inflammatory cell recruitment; immune function regulation; model; three-dimension; 3-D;

Homo sapiens

12-AUG-1999

04-FEB-1999; 99WO-IB00367

11-SEP-1998; 06-FEB-1998; 98US-0099994. 98US-0073972.

(ILEX-) ILEXUS PTY LTD

Baell JB, Maxwell KF, Epa V, Garrett TPJ, Hog, McCarthy TD, McKenzie Hogarth PM, zie IFC, Pi Pietersz GA; Matthews BR

WPI; 1999-539978/45.

Three-dimensional structures and models of Fc receptors, useful computer-assisted drug design $% \left\{ 1\right\} =\left\{ 1$

Claim 11; Page 316-317; 326pp; English

reduce IgG- or IgE-mediated tissue damage, e.g. IgG- or IgE-mediated hypersensitivity, recruitment of inflammatory cells or release of inflammatory modulators. The therapeutic compositions can also be used to treat disease in animals. The knowledge of the 3-D structure of FC-gamma RIa and models of other FCR provides a means for designing and producing compounds that regulate immune function and inflammation in an animal, including humans (i.e. structure based drug design). For example, chemical compounds can be designed to block binding of immunoglobulin to an FC recentor protein main. This invention describes a novel model of an Fc receptor (FCR) protein representing a three-dimensional (3-D) structure that substantially conforms to the specified atomic coordinates. Computer model images of an Fc receptor protein using various computer programs and models. The products of the invention have anti-inflammatory and immunoprotective activity. This sequence represents the human sFC-gammaRI protein oligonucleotides, peptides, peptidomimetics and small organic moleculous computer-assisted methods of drug design. Therapeutic compositions that inhibit the activity of FC-gammaR or FC-epsilonR can be used to the FcR can be used to design bioactive chemical fragment described in the method of the pn bioactive chemical compounds, e.g.
peptidomimetics and small organic molecules. ç

Sequence

Query Match Best Local (Matches Local Similarity 1 GNLVTLSCETKL 12; Conservative 12 100.0%; 0; Score 61; DB 20; Pred. No. 0.0073; Length 261; Indels 0; Gaps

0;

RESULT R22549

QY

밁

186 gnlvtlscetkl 197

R22549 standard; Protein; 344 AA

21-MAY-1992 (first entry)

Human macrophage-specific FcRI receptor encoded by clone p98

Rapid immunoselection cloning technique immunodiagnosis; high affinity receptor technique; cell surface antigen;

Homo sapiens

W09201049-A

XX PX XX

23-JAN-1992.

15-JUL-1990; 90WO-US04986

13-JUL-1990; 90US-0553759

PAX PR

(GEHO-) GEN HOSPITAL CORP

Seed B, Aruffo A, Amiot M;

N-PSDB; 1992-056864/07 DB; Q21179.

New CD53 cell surface antigen and DNA encoding immuno-therapy and diagnosis of haematopoietic $% \left(1\right) =\left\{ 1\right\}$ it - for
neoplasms,

CCCCXXXX PTX TXXX

Example 10; Page 94a; 160pp; English

This amino acid sequence was predicted from the cDNA sequence of cDNA clone 998/X2. It differs from the sequence predicted from clone 9135 (see 021178) at position 25; 9135 encodes a Ser residue while clone 998/X2 predicts a Thr residue. At position 58, 990Ser residue 58, p90

B 20

0

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RESULT
R91439
ID R9
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Best Local S
Matches 12
immunoselection cioning method. The for the FC portion of IgG, normally located on cell surfaces of macrophages. 2 Other cDNA clones, pl35 (T14717) and p90 (T14719), coded for a variants of the FCRI sequence; the C-terminal sequence coded for a variant of the FCRI sequence with those of the pl35
                                                                                                                                                                                                                          Cloning of cDNA encoding cell surface antigen of diagnostic and therapeutic proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-DEC-1992;
25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cys residues distributed among three C2 set Ig-related domains. A fusion protein of FCRI and a receptor ligand will be helpful to increase the potency of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane-spanning domain, and a short, highly charged cytoplasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and six
                                                                                           cDNA clone, p98 (T14718), obtd. from a cDNA library using an immunoselection cloning method. FcRI is a high affinity reco
                                                                                                                                                                                                                                                                                          N-PSDB; T14718
                                                                                                                                                                                                                                                                                                                                                            Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           09-APR-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cell surface antigen;
therapy; diagnosis; ve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human FcRI (cDNA clone p98 product).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         R91439;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  R91439 standard; Protein; 344 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                 The amino acid sequence (R91439) of human FcRI was detd. from a
                                                                                                                                                                            Example 10; Column 55-56; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  18-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-JUL-1989
13-JUL-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US5506126-A
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                                                                                                                                                                                                                                                                                                                                                                                                     (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             205
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                                                                                                                                                                                                                                                                                                                                                            Seed B;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                              89US-0379076.
90US-0553759.
93US-0139273.
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88US-0160416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               88US-0160416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     vector;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 61; 100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cloning; immunoselection; immunotherapy;
ector; FcRI; Fc receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "amino acid
translated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               158 is Leu
1 product"
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0.0097;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 in clone pl35
                                                                                                                                                                                                                                                 useful for isolation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 344;
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Y96183
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   macropusys, recall panning; immunodagnosis; diagnosis; immunotherapy; immune disorder; infection; asthma; immune-complex complex of the control of the contr
                                        Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            205
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gnlvtlscetkl 216
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213
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136
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123
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122
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117
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139 ·
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A fusion protein in increasing the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 344;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                disease;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                therapy;
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В

205 gnlvtlscetkl 216

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01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
23-MAR-1990;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Misc-difference
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                                          sclerosis. FCRI is a high affinity receptor for the FC polific normally located on the cell surfaces of macrophages ability to interfere with such bonding, or to cause it to cause
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Isolated nucleic acid molecule encoding the CD19 cell surface antigen, useful for immunodiagnosis and immunotherapy of immune-mediated infections or disorders, e.g. asthma, immune-complex disease, parasiting the complex disease, parasiting the complex disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Stamenkovic I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-OCT-1998;
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surfaces other than macrophages, protein of FCRI and a receptor 1
                                                                                                                                                                                                                                                                                    from a CDNA Library expression of a CSA nucleic acids. The method is the invention designed to isolate CSA nucleic acids. The method is based upon transient expression of a CSA in eukaryotic cells and based upon transient expression the antigen by adhesion to
                                                                                                                                                                                                                                                                                                                                                                                  The present sequence is that of a human macrophage specific FCRI, as deduced from cDNA clone p135 (see A50631), which was isolated
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 10; Column 53-55;
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                                                                                                                                        animals, including humans.
                                                                                                                                                                 immunotherapeutic applications, including the diagnimmunotherapeutic applications, including the diagnimunotherapeutic applications, diseases,
                                                                                                                                                                                                                   proteins they encode, are useful for immunodiagnostic and
                                                                                                                                                                                                                                       CSA nucleic acids isolated by the method of the invention, and the
                                                                                                                                                                                                                                                                ohysical selection of cells expressing the antigen by adhesion to (panning on) an antibody-coated substrate such as a culture dish.
                                                                                            mmune-complex disease, amyloidosis, parasitic diseases or multiple sclerosis. FcRI is a high affinity receptor for the Fc portion of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2000-586382/55
                                                                                                                                                                                                                                                                                                                                                              uced from cDNA clone p135 (see A50631), which was isolated cDNA library expressed in COS cells using a novel method of
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306
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88US-0160416
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216
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305
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220
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338
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332
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        75pp;
                                                                                                                                             These disorders include asthma,
bonding, or to cause it to occur on is useful in therapy. A fusion ligand will be helpful to increase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       by
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                                                                                                                                                                                             the diagnosis and
                                                                                                                                                                    and disorders in
                                                                                                                                                                                                                                                                                                                                             The method is
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Matches

GNLVTLSCETKL 12

Query Match 'Best Local Similarity

100.0%;

Score 61; DB 21; Pred. No. 0.0097; 0; Mismatches (

0;

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Gaps

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21-MAY-1992

(first entry)

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DB 21;

Length 344; Indels

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Sequence

344 AA; of

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X D X C X U
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                                                                                                                    RESULT
                                                                                                                                                                                                                                         Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 12; Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                typical features of a type I integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane-spanning domain, and a short, highly charged cytoplasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and six Cys residues distributed among three C2 set Ig-related domains. A fusion protein of FCRI and a receptor ligand will be helpful to increase the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This amino acid sequence was predicted from the cDNA sequence of cDNA clone pl35. It differs from the sequence predicted from clones p98/X2 and p90 (see Q21179 and Q21180, respectively) at position 25, pl35 encodes a Ser residue and the other two clones predict a Thr residue. At position 58, pl35 predicts Leu and p90 predicts Val. Sequences predicted from all 3 clones show the predicts Val. Sequences predicted from all 3 clones show the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New CD53 cell surface antigen and DNA encoding immuno-therapy and diagnosis of haematopoietic % \left( 1\right) =\left\{ 1\right\} 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rapid immunoselection cloning technique; cell surface antigen; immunodiagnosis; high affinity receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human macrophage-specific FcRI receptor encoded by clone p135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  R20811 standard; Protein;
                                                                                                                                                                                                                                                                                                                                  Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-JUL-1990;
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                                                                                                                                                                                                                                                                                                                                                                 potency of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 10; Page 94a; 160pp; English:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (GEHO-) GEN HOSPITAL CORP
                                                 R22550;
                                                                                   R22550 standard;
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                                                                                                                                                                                                                                                                                                                                    374 AA;
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                                                                                     Protein; 374 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      374 AA.
                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                               Score 61; DB 13
Pred. No. 0.011;
                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                 DB 13;
                                                                                                                                                                                                                                                0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          it -
                                                                                                                                                                                                                                                                                 Length 374;
                                                                                                                                                                                                                                                                                                                                                                                                           A fusion protein
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Best Local Similarity
Matches 12; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             domain, and a short, highly charged cytoplasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and six Cys residues distributed among three C2 set Ig-related domains. A fusion protein of FCRI and a receptor ligand will be helpful to increase the potency of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           and clone p90 predicts a Thr residue. At position 58, p90 predicts val while clones p135 and p98/X2 (see Q21179) both predict Leu. Sequences predicted from all 3 clones show the typical features of a type I integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane-spanning
                                                                                                        Cell surface antigen; cloning; immunoselection; immunotherapy;
therapy; diagnosis; vector; FcRI; Fc receptor.
                                                                                                                                                                          30-OCT-1996
                                                                                                                                                                                                                              R91438 standard; Protein; 374 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Seed B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Rapid immunoselection cloning technique; cell surface antigen; immunodiagnosis; high affinity receptor.
Misc-difference
                                                                               Homo sapiens
                                                                                                                                                Human FcRI (cDNA clone pl35 product).
                                                                                                                                                                                                     R91438
                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 10; Page 94a; 160pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New CD53 cell surface antigen and DNA encoding immuno-therapy and diagnosis of haematopoietic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       N-PSDB; Q21180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1992-056864/07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23-JAN-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               W09201049-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                         Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human macrophage-specific FcRI receptor encoded by clone p90
                                                                                                                                                                                                                                                                                              205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (GEHO-) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                        1 GNLVTLSCETKL 12
                                                                                                                                                                                                                                                        8
                                                                                                                                                                                                                                                                                              gnlvtlscetkl 216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                    374 AA;
                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                90us-0553759
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           It differs from the sequence predicted from 221178) at position 25; pl35 encodes a Ser residue
  58
                                        Location/Qualifiers 25
                             /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Amiot M;
                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 61; DB 13
100.0%; Pred. No. 0.011;
tive 0; Mismatches
             "amino acid 25 is Thr in
translated products"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    English
                                                                                                                                                                                                                                                                                                                                                                          DB 13;
                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             neoplasms,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           it - for
                          clone p90 and p98
                                                                                                                                                                                                                                                                                                                                                                         Length 374;
                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              etc.
                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         XXCCCCCCCXXX
                                                                                                                                                                                                                                                                                              W00859
                                                                                                                                                                                                                                                                                                             RESULT
                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
18-OCT-1993;
01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
                                                                               09-APR-1996
                                                                                                         US5506126-A
                                                                                                                                                           Cell surface antigen; cloning; immunoselection; immunotherapy; therapy; diagnosis; vector; FCRI; Fc receptor.
                                                                                                                                                                                                  Human FcRI (cDNA clone p90 product).
                                                                                                                                                                                                                              30-OCT-1996
                                                                                                                                                                                                                                                                                W00859 standard; Protein; 374 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 10; Column 55-56; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    N-PSDB; T14717.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1996-200279/20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         09-APR-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ŪS5506126-A
                                                    25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GEHO ) GEN HOSPITAL CORP.
                                                                                                                                                                                                                                                                                                                                                205 gnlvtlscetkl 216
                                                                                                                                                                                                                                                                                                                                                              1 GNLVTLSCETKL 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Seed B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             92US-0983647.
88US-0160416.
89US-0379076.
99US-0553759.
93US-0139273.
             92US-0983647.
88US-0160416.
                                                      88US-0160416
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                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%;
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The amino acid sequence (R91438) of human FCRI was detd. from a cDNA clone, p135 (T14717), obtd. from a cDNA library using an immunoselection cloning method. FCRI is a high affinity receptor for the Fc portion of IgG, normally located on cell surfaces of macrophages. Another isolated cDNA clone, p90 (T14719), coded for a variant (W00859) of the FCRI sequence, and a third clone, p96 (T14718), coded for an FCRI (R91439) having a different C-terminal sequence. A fusion protein of FCRI and a receptor ligand will be helpful in increasing the potency of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cloning of cDNA encoding cell surface antigen - useful of diagnostic and therapeutic proteins % \left( 1\right) =\left\{ 1\right\} =\left\{ 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "amino acid 58 is Val in p90 clone translated product"
                                                                                                                                                                                     Score 61;
Pred. No.
Mismatches
                                                                                                                                                                                     0.011;
                                                                                                                                                                                                                                                                                                                                                                                               DB 17;
0
                                                                                                                                                                                                                                                                                                                                                                           Length 374;
                    Indels
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0;
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Gaps
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Best Local Similarity
                                                                                                                                                                           /note=
Misc-difference 45
                                                                                                                                                                                                                                                                                                                                                                                                    07-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The amino acid sequence (W00859) of human FCRI was detd. from a CDNA clone, p90 (T14719), obtd. from a cDNA library using an immunoselection cloning method. FCRI is a high affinity receptor for the FC portion of IgG, normally located on cell surfaces of macrophages. Other CDNA clones (see also T14717-18) coded for the FC DNA clones (see also T14717-18) coded for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cloning of cDNA encoding cell surface antigen of diagnostic and therapeutic proteins
         Misc-difference
                                Misc-difference
                                                           Misc-difference
                                                                                                              Misc-difference
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                                                                                                                                                                Misc-difference
                                                                                                                                                                                                                    Misc-difference
                                                                                                                                                                                                                                             Misc-difference
                                                                                                                                                                                                                                                                                                                                     cloning.
                                                                                                                                                                                                                                                                                                                                                                        Human Fc receptor I
                                                                                                                                                                                                                                                                                                                                                                                                                            W80448;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   W80448 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           variants (R91438-39) of the FCRI sequence. A fusion protein FCRI and a receptor ligand will be helpful in increasing the potency of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 10; Column 55-56; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1996-200279/20
N-PSDB; T14719.
                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                Fc receptor I; FcRI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13-JUL-1990;
18-OCT-1993;
                                                                                                                                                                                                                                                                       Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (GEHO ) GEN HOSPITAL CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 GNLVTLSCETKL 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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93US-0139273
                                                                                                                /note=
85
        /note=
                                                                                     /note=
99
                                                                                                                                                                /note=
                                                            103
                                                                                                                                        /note=
77
                                                                                                                                                                                                                                             /note=
23
                                                                                                                                                                                                                                                                          Location/Qualifiers
                                             'note=
                                                                       'note=
                                                                                                                                                                                                                             'note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%;
                                                                                                                                                                                                                                                                                                                                               cell surface antigen; lymphocyte; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          in therapy.
                                                                                                                                                                                                      "encoded
                  encoded by
                                         "encoded by CCC"
                                                                                                                                                  "encoded by CCC"
                                                                                                                                                                           "encoded
                                                                                                                                                                                                                                "encoded
                                                                                                                                                                                                                                                        "encoded by TGG"
                                                                       "encoded
                                                                                                 "encoded
                                                                                                                           "encoded
                                                                                                                                                                                                                                                                                                                                                                                                                                                    374 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 61;
Pred. No.
                                                                                                                        ΛĠ
                                                                    by CAA"
                                                                                             by TCC
                                                                                                                                                                           by CTG"
                                                                                                                                                                                                      by GAC"
                                                                                                                                                                                                                               by CTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                     GGC"
                                                                                                                           AAT"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.011;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 17; Length 374;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A fusion protein of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     useful for isolation
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RESULT W97833

11

AC XX

W97833 standard; Protein; 374 AA.

W97833;

В

205 gnlvtlscetkl 216

1 GNLVTLSCETKL 12

Qy

Query Match 100.0%; Best Local Similarity 100.0%; Matches 12; Conservative

0

Mismatches

Score 61; DB 19 Pred. No. 0.011;

19; 0;

Length 374; Indels

of

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Gaps

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Sequence

374

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88888888888888888888888888888888888
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13-JUL-1990;
21-MAY-1997;
                                                                                                                                                    transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V63442-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated genes can be expressed in a prokaryotic or eukaryotic host cells to produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V63444) which allow
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note=
Misc-difference 256
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note=
Misc-difference_176
                                                             the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and immunotherapeutic applications, vincluding the diagnosis and treatment of immune-mediated infections, diseases, and disorders
                                                                                                                                                                                                                                                                                     in eukaryotic cells and selection of cells expressing the antigen by adhesion to an antibody-coated substrate. The method is useful for the isolation and cloning of any protein which can be expressed and
                                                                                                                                                                                                                                                                                                                                    were also obtained. A novel method for cloning cDNAs from mammalian expression libraries is based on transient expression of an antigen by in eukaryotic cells and selection of cells expressing the antigen by
                                                                                                                                                                                                                                                                                                                                                                                 were also obtained.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This is the amino acid sequence of human Fc receptor I (FcRI), deduced from cDNA clone pl35 (see V63456) isolated using a rap
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 10; Column 53-54; 75pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              03-NOV-1998
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                                                                                                                                                                                                                                                                                                                                                                              from a single patient undergoing extracorporeal interleukin-2 induction therapy. Clones p90 (see x07372) and p98 (see x07373) were also obtained. A novel method for cloning cDNAs from mamma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            human lymphocyte antigenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   N-PSDB; V63456
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-DEC-1992;
25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US5830731-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        immunoselection cloning method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1998-609251/51
                                                                                                                                                                                                                                                                                                                                                                                                                                                     The cDNA library was constructed
                                            including humans.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Seed B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   89US-0379076
90US-0553759
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              97us-0861205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                92US-0983647
88US-0160416
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171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "encoded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "encoded by AAC"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          from a cDNA library expressed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             γď
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GTC"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATG"
                                                                                                                                                                                                                                                                                                                                                                                                                                                     from polyA RNA of cells
                                                                    disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               rapid
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25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note= "encoded Misc-difference 99
                                             This is the amino acid sequence of human Fc receptor I (FcRI), as deduced from cDNA clone p90 (see X07372) isolated using a rapid immunoselection cloning method from a cDNA library expressed in COS cells. The cDNA library was constructed from polyA RNA of cells
                                                                                                                                         New cloning vector and polylinker - based on existing sequences fefficient cloning and expression of mammalian cDNA(s), especially human lymphocyte antigenic sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note=
Misc-difference 171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note=
Misc-difference 85.
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N-PSDB; X07372.
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         from a single patient undergoing extracorporeal induction therapy. Clones pl35 (see V63456) and were also obtained. A novel method for cloning
                                                                                                                Example 10; Column 53-54; 75pp; English.
                                                                                                                                                                                                                                     Aruffo A,
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is based on transient expression

of an antigen

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V63442-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated
                                                                                                                                                                                                                                                                                                  Misc-difference
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                                  Misc-difference
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Best Local Similarity
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               Macrophage: FCRI; cell surface antigen; human; immunoselection; panning; immunodiagnosis; diagnosis; immunotherapy; gene therapy; immune disorder; infection; asthma; immune-complex disease; amyloidosis; multiple sclerosis; parasitic disease; Fc receptor.
                                                                                                                     Human macrophage-specific FcRI.
                                                                                                                                                                                                                                              Y96134 standard; Protein;
                                                                                                                                                                19-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New cloning vector and polylinker - based on efficient cloning and expression of mammalian human lymphocyte antigenic sequences
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2000-586382/55

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RESULT 14
Y96226
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                                                                                                                                                                                                                  Human; high affinity Fc receptor; FcgammaRI; immunoglobulin; infection; immune response; CD64; monocyte; macrophage; neutrophil; eosinophil; HIV: 19G; immunosuppressive; antirheumatic; cytostatic; antiinflammatory; AIDS; SLE; multiple myeloma; rheumatoid arthritis;
                                                          03-DEC-1998;
                                                                                                                   07-JUN-2000
                                                                                                                                              EP1006183-A1
                                                                                                                                                                                                                                                                                                                                                       Y96226
WPI; 2000-367968/32
                                                                                     03-DEC-1998;
                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                     systemic lupus
                                                                                                                                                                                                                                                                                           Human high affinity Fc receptor, FcgammaRI
                                                                                                                                                                                                                                                                                                                                                                                   Y96226 standard; Protein; 374 AA.
                           (PLAC ) MAX PLANCK GES
                                                                                                                                                                                                                                                                                                                                                                                                                                                           205
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                       erythematosus; tumour
                                                          98EP-0122969.
                                                                                       98EP-0122969
                             FOERDERUNG WISSENSCHAFTEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 61; DB 21
Pred. No. 0.011;
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receptor consist only of the extracellular portion of the receptor and are not glycosylated i.e. they do not have transmembrane domains or signal peptides. The recombinant proteins my be used in immunoassays to determine the immune status of patients with chronic diseases of the immune system, e.g. AIDS, systemic lupus erythematosus (SLE), multiple myeloma (MM), or rheumatoid arthritis. In addition, pharmaceutical
                                                                                                                                                                                                                                                                                             monocytes and macrophages, but expression can also be induced on neutrophils and eosinophils. Upon Fc receptor activation, immune effecto pathways are activated, leading to immune response. The present sequence
                                                                                                                                                                                                                                                                                                                                                                    role in defending the body against infections. First, pathogens are opsonised by serum immunoglobulins. The resulting complex then binds cells expressing Fc receptors. FcgammaRI molecules are expressed by
                                                 prevent autoimmune diseases, allergies or tumours, especially AIDS rheumatoid arthritis or MM.
                                                                                             myeloma (MM), or rheumatoid arthritis. In addition, pharmaceutical compositions containing recombinant proteins may be used to treat or
                                                                                                                                                                                                                                                                      may be modified to produce recombinant versions.
                                                                                                                                                                                                                                                                                                                                                                                                                     role in defending the body against infections. First, pathogens are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel Fc receptor lacking transmembrane domains, a signal peptide, and glycosylation, useful for diagnosing and treating immune disorders and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sequence is the human high affinity Fc receptor,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             26-28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     60pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          English
                                                                                                                                                                                                                                                                           The recombinant Fc
                                                                                                                                                                                                                                                                                                                             immune effector
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FcgammaRI
                                                                                                                                                                                                                                                                                                                                                                                                     6
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The present sequence is that of a human macrophage specific FCRI, as deduced from cDNA clone p135 (see A50592), which was isolated from a cDNA library expressed in COS cells using a novel method of the invention designed to isolate CSA nucleic acids. The method is based upon transient expression of a CSA in eukaryotic cells and

Example 10; Column 53-55; 75pp; English

physical selection of cells expressing the antigen by adhesion to (panning on) an antibody-coated substrate such as a culture dish. CSA nucleic acids isolated by the method of the invention, and the

and the

is

proteins they encode, are useful for immunodiagnostic and

immunotherapeutic applications, including the diagnosis and disorders in

Isolated nucleic acid molecule encoding the CD19 cell surface antigen, useful for immunodiagnosis and immunotherapy of immune-mediated infections or disorders, e.g. asthma, immune-complex disease, parasitic

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Matches
                                     Query Match
Best Local
1 GNLVTLSCETKL 12
                              Similarity
12; Conserv
                               Conservative
                                     100.0%;
                               0
                                      Score 61;
Pred. No.
                               Mismatches
                                      0.011;
                                             DB 21;
                                             Length 374;
                               Indels
                               0
                               Gaps
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0;

Query Match Best Local

Local

Similarity

100.0%;

21;

Length 374;

밁 QΥ Sequence

374 AA;

ability to interfere with such bonding, or to cause it to occur on surfaces other than macrophages, is useful in therapy. A fusion protein of FCRI and a receptor ligand will be helpful to increase

lmmune-complex disease, amyloidosis, parasitic diseases or multiple sclerosis. FcRI is a high affinity receptor for the Fc portion of

These disorders include asthma,

including humans.

normally located on the cell surfaces of macrophages.

The

Sequence

374

AA;

the potencies of antibodies in therapy.

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RESULT
Human cancer associated protein sequence SEQ ID NO:1128
                                          08-FEB-2001
                                                                                                                       B43683 standard; Protein; 399 AA.
                                                                                                                                                               15
                                       (first entry)
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neurological disease; vasotropić; antipsoriatic; antiangiogenic; gene therapy; inflammation; immune disorder; haematopoletic cell disorder; autoimmune disorder; allergic reaction; graft versus host disease; organ rejection; haemostatic; thrombolytic; cardiovascular disorder; infection; antidiabetic; antiasthmatic; antirheumatic; antiarthritic; antiviral; diagnosis; dermatological; antiinflammatory; antithyroid; antiallergic; antibacterial; cardiant; cancer associated gene; cancer antigen; detection; cytostatic; neuroprotective; thrombolytic; proliferative; vulnerary; coagulant; nootropic 1mmunomodulator; cancer;

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Rosen CA;
                                       08-MAR-2000; 2000WO-US05882
                                                    21-SEP-2000
                                                                              Homo sapiens
            (HUMA-) HUMAN GENOME SCI
                          12-MAR-1999;
                                                                 WO200055350-A1
Ruben SM;
                          99US-0124270.
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WPI; 2000-587533/55

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N-PSDB; C77892.
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Novel isolated nucleic acids comprising sequences encoding peptides useful for treating or diagnosing e.g. cancer - $\,$

Claim 11; Page 1739-1740; 2352pp; English.

antithyroid; antiallergic; antibacterial; antiviral; dermatological; consurprotective; cardiant; thrombolytic; coagulant; nootropic; vasotropic; antipsoriatic and antiangiogenic. The polynucleotides and conditions and diagnosing pathological conditions. Polynucleotides, conditions and be used to treat immune disorders by activating or inhibiting the proliferation, differentiation or mobilisation of immune conhibiting the proliferation, differentiation or mobilisation of immune conhibiting the proliferation, differentiation or mobilisation of immune collegic reactions, graft versus host disease and organ rejection, condulate haemostatic or thrombolytic activity, modulate inflammation, concers, cardiovascular disorders, neurological disease and bacterial or viral infections. The peptides, nucleotides, antibodies, agonists and content and content sequences used in drug screens. C78449 to C78457 and sequences used in the exemplification of the present incomplete. C77607 to C78448 encode the human cancer associated proteins given in B43398 to B44239. The proteins can have activities based on the tissues and cells the genes are expressed in Example of activities include: antiasthmatic; antirheumatic; antiarthritic; antiinflammatory; cytostatic; proliferative; vulnerary; immunomodulator; antidiabetic; invention.

Sequence 399 AA;

Query Match 100.0%; Score 61; DB 21; Length 399; Best Local Similarity 100.0%; Pred. No. 0.011; Matches 12; Conservative 0; Mismatches 0; Indels Indels 0; Gaps 0;

B 1 GNLVTLSCETKL 12 ||||||||||| | 230 gnlvtlscetkl 241

Search completed: June 4, 2001, 12:13:27 Job time: 200 sec

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         of hits satisfying chosen parameters:
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3: /cgn2_6/ptodata/2,
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Query Match
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78.78;

Score 48;

DB 2;

Length 91;

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US-08-466-860-7
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JS-08-297-395-10
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Matches 8; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ARLIER APPLICATION NUMBER: 08/059,189
ARLIER FILING DATE: 1993-05-06
ARLIER APPLICATION NUMBER: 07/502,559
ARLIER FILING DATE: 1990-03-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        JRRENT APPLICATION NUMBER: UPPRENT FILING DATE: 1994-08-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ARLIER APPLICATION NUMBER: 07/065,734
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TLE OF INVENTION:
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/466,860
                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LE REFERENCE:
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RLIER FILING DATE: 1988-06-24
                                                                                                                                                                                                                                          ORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                     PPLICANT:
                                                   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                              ITLE OF INVENTION: RESULTING FI
                                                                                                                                                                                                                                                                                                                  ITLE OF INVENTION:
                                                                                                                                                                  CITY: SAN DIEGO
STATE: CALIFORNI
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                                                                                                                                    COUNTRY: 1
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|7 GSLVTLNCETNL 28
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                                                                                                                                               CALIFORNIA: UNITED STATES
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                                                                                                                                                                                                    E: CAMPBELL AND FLORES 4370 LA JOLLA VILLAGE.
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                                                                                                                                                                                                                                                                                                                                                                       HOWELL, MARK D
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                                                                                                                                                                                                                                                                                             VACCINATION AND METHODS AGAINST DISEASES RESULTING FROM PATHOGENIC RESPONSES BY SPECIFIC T CELL
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Best Local :
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                                                                                                               TELEFAX: 619-535-8949
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ENERAL INFORMATION:
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ATTORNEY/AGENT INFORMATION:
NAME: CAMPBELL, CATHRYN
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PRIOR APPLICATION DATA:
     MOLECULE TYPE: FRAGMENT TYPE:
                                                                           SEQUENCE CHARACTERISTICS: LENGTH: 113 amino acid
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                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION: NAME: CAMPBELL, CATHRYN
                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/0 FILING DATE: 06-JUN-1995
                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                       REGISTRATION NUMBER: 31. REFERENCE/DOCKET NUMBER:
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STATE: CALIFORNIA
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                                                                                                                                                 TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                     SOFTWARE:
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                                             TOPOLOGY:
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4370 LA JOLLA VILLAGE DRIVE, SUITE 700
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N-terminal
       peptide
N-terminal
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RESULTING FROM PATHOGENIC RESPONSES BY SPECIFIC T CELL
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Pred. No. 13;
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Length 113; Indels

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Best Local Similarity
Matches 8; Conser
              Sequence 14, Appli
Patent No. 5686600
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GENERAL INFORMATION:
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NFORMATION FOR SEQ ID NO:
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LENGTH: 113 amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                              NAME: CAMPBELL, CATHRYN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
PRIOR APPLICATION DATA:
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SURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: CATTO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                JMBER OF SEQUENCES:
                                                                                                  35 GQNVTLSCEQNL 46
                                                                                                                                                                                                                                                                      TOPOLOGY:
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                                                                                                                           GNLVTLSCETKL 12
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                                                                                                                                                                                                                                                                                  amino acid
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                          Application US/08442542
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                                                                                                                                                                                                                                                                                               113 amino acids
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CARLO, DENNIS J.

VENTION: VACCINATION AND METHODS AGAINST DISEASES

VENTION: RESULTING FROM PATHOGENIC RESPONSES BY SPECIFIC T CELL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             UNITED STATES
                                                                                                                                                       Conservative
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N-terminal
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Pred. No.
                                                                                                                                                                     Score 37;
Pred. No.
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Query Match
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                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                tent No.
                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
COMPUTER: IBM PC com
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
            SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                       TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                         CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 0 FILING DATE: 28-JUN-1994 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION DATA:
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                     COMPUTER: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
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SOFTWARE: PatentI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
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                                                                                                                                              ITY: Hawthorne
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SYSTEM: PC-DOS/MS-DOS
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70.0%;
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US/08/765,469
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Pred. No.
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Best Local Similarity
Thes 7; Conserv
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US-08-765-469-14
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GENERAL INFORMATION:
APPLICANT: Presid
                                                             TELEFAX: 207-363-0528
NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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INFORMATION FOR SEQ ID NO: '14:
SEQUENCE CHARACTERISTICS:
                                                                                                                 TELECOMMUNICATION INFORMATION: TELEPHONE: 207-363-0558
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: President and Fellows of Harvard College TITLE OF INVENTION: SINGLE CHAIN T-CELL RECEPTOR NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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NAME: Spruill, W. Murray
REGISTRATION NUMBER: 32,
                                                                                                                                                TTTORNEY/AGENT INFORMATION:
NAME: Farrell, Kevin M;
REGISTRATION NUMBER: 35,505
REFERENCE/DOCKET NUMBER: HU
                                                                                                                                                                                                                   RIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/349.915
FILING DATE: -- O6-DEC-1994
CLASSIFICATION:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 919-541-8615
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APPLICATION NUMBER:
                   TOPOLOGY:
                               TYPE: amino acid
                                                                                                                                                                                                                                                                                        CLASSIFICATION:
                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: PCT/US95/15696
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                                               ENGTH:
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                                               391 amino acids
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Pred. No. 14;
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                                                         Matches
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LIBRARY: GenBar
CLONE: 969078
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                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  quence 3, Application US/08896005
tent No. 5854023
                                                                                                                                                                                                                                                                              TELEPHONE: 415-855-0555
                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF
                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/8 FILING DATE: Filed Herewith PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: FastSEQ for Windows Version 2.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS
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403 GRLVNLSCST 412
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                                                                                                                                                                                                           TYPE: amino acid
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                           1 GNLVTLSCET 10
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Corley, Neil C
                                                                                                                                                     GenBank
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70.0%;
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                                                                   Score 36; DB Pred. No. 91;
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                                                       Mismatches
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RESULT 10 US-08-037-816A-20

Sequence 20,

Application US/08037816A

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Patent No. 5869624
GENERAL INFORMATION:
APPLICANT: Hasel,
APPLICANT: Maddon,

Hasel, Karl W. Maddon, Paul J.

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Query Match
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Best Local Similarity
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SEQUENCE CHARACTERISTICS:
LENGTH: 479 amino acids
TYPE: amino acid
            SOFTWARE: PatentIn Release #1.24 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/037,816A FILING DATE: 26-MAR-1993
                                                                            COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPUTER: DOS MS DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFERRICE/DOCKET NUMBER: 41190/JPW/AJM
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 977-9550
TELEPHONE: (212) 977-9550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Relaction CATA:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                    ORRESPONDENCE ADDRESS:
ADDRESSEE: Cooper & Dunham
                                                                                                                                                                                                                                                                                         FITLE OF INVENTION:
FITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                           APPLICANT:
                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
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                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
                                                                                                                                                                                     CITY:
STATE:
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CLASSIFICATION:
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CLASSIFICATION:
                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                      STREET: - 30 Rockefeller Plaza
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                                                                                                                                                  10112
                                                                                                                                                                                New York
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                                                                                                                                                                                                                                                                                                                                                           Hasel, Karl W.
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Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 41190/JPW/AJM
TELECOMMUNICATION INFORMATION:
TELEPHAX: (212) 977-9550
TELEFAX: (212) 664-0525
TELEEX: 422523 COOPUI
INFORMATION FOR SEQ ID NO: 24:
SEQUENCE CHARACTERISTICS:
LENGTH: 479 amino acids
                                               INFORMATION FOR SEQ ID NO: 20: SEQUENCE CHARACTERISTICS: LENGTH: 479 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SENERAL INFORMATION:
                                                                                                                                     NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 411
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 977-9550
                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PM PC POS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.24
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/037,816
FILING DATE: 26-MAR-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION: NAME: White, John P.
MOLECULE TYPE:
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                                                                                                                     TELEFAX:
                                                                                                                                                                                                                                                                                                          FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: New York
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET:
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                 TOPOLOGY:
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                              amino acid
                                                                                                      ONE: (212) 977-95
4: (212) 664-0525
422523 COOPUI
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Maddon, Paul J.
VENTION: HIV-1 VACCINES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           USA
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Pred. No. 1.3e+02;
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57.48;

Score 35; Pred. No.

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RESULT 14
US-08-037-816A-16
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US-08-530-146-24
                                                                     Sequence 16, Application US/08037816A Patent No. 5869624
                                                                    Patent No.
                                                                                                                                                                                                                                                                                              TOPOLOGY: 11
MOLECULE TYPE:
-08-530-146-24
                                                       GENERAL INFORMATION:
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equence 24, AFF-
No. 5886163
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                        APPLICANT:
                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS LENGTH: 479 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.24
            TTLE OF
                                                                                                                                                          380 GNTITLPCRIK 390
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                             ELECOMMUNICATION INFORMATION: TELEPHONE: (212) 977-9550
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER:
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THE MADD PAUL J. INVENTION: HIV-1 VACCINES, ANTIBODY COMPOSITIONS RELATED
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                                        Hasel, Karl W.
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                                                                                                                                                                                                                    Score 35; DB 2;
Pred. No. 1.3e+0;
1; Mismatches
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                                                                                                                                                                                                                                                Length 479;
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Best Local Similarity 54.9
Matches 6; Conservative
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                                                                                                                                                                                                                                                                                                                                                                        atent No. 5869624
GENERAL INFORMATION:
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                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                              CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/037,816A
                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 507 amino acids
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MEDIUM TYPE: Floppy
COMPUTER: IBM PC com
                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                APPLICANT: Maddon, ITLE OF INVENTION:
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TTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                   ITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              408 GNTITLPCRIK 418
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: (212) 977-9550
                                                                             COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Palanta
                                FILING DATE:
                                                                                                                                                                                                CITY: New York
STATE: New York
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               CLASSIFICATION:
                                                                                                                                                                              COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: FILING DATE: 26-MAI CLASSIFICATION: 43
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: UZIP: 10112
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30 Rockefeller Plaza
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                                                                                              IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 35; DB 2;
Pred. No. 1.4e+02
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REGISTRATION NUMBER: 28,678
REGISTRATION NUMBER: 41190/JPW/AJM
REFERENCE/DOCKET NUMBER: 41190/JPW/AJM
RELECOMMUNICATION IMFORMATION:
TELECOMMUNICATION IMFORMATION:
TELEPHONE: (212) 977-9550
TELETAX: (212) 664-0525
TELEX: 42253 COOPUI
INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
LENGTH: 507 amino acids
TYPE: amino acids
TYPE: amino acids
TYPE: molecula Type: protein
US-08-037-816A-28

Ouery Match
Best Local Similarity 54.5%; Score 35; DB 2; Length 507;
Best Local Similarity 54.5%; Pred. No: 1.4e+02;
Matches 6; Conservative 1; Mismatches 4; Indels 0; Gaps

Outy Match
Ouery Match
Best Local Similarity 54.5%; Pred. No: 1.4e+02;
Matches 6; Conservative 1; Mismatches 4; Indels 0; Gaps

Ouery Match
Ouery Match
Best Local Similarity 54.5%; Pred. No: 1.4e+02;
Matches 6; Conservative 1; Mismatches 4; Indels 0; Gaps

Ouery Match
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Search completed: June 4, 2001, 12:14:29

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A;Title: Isolation and expression of functional high-affinity Fc receptor complementa A;Reference number: A41357; MUID:89100284
A;Accession: A41357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Species: Homo sapiens (man)
C;Date: 03-Apr-1992 #sequence_revision 03-Apr-1992 #text_change 21-Jan-2000
C;Accession: A41357; S03019
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A41357
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A; Residues: 1-344 < ALI
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ALIGNMENTS

	A; Restudes: 1-34 (ALLI) A; Cross-references: GB:X14355; GB:M21090; NID:g31333; PIDN:CAA32536.1; PID:g31334 A; Closs-references: GB:X14355; GB:M21090; NID:g31333; PIDN:CAA32536.1; PID:g31334 A; Reference Acids Res. 16, 11824, 1988 A; Pittle: Nucleotide sequence of three cDNAs for the human high affinity Fc receptor (A; Reference number: S03018; MUID:89098339 A; Reference number: S03018; MUID:89098339 A; Accession: S03019 A; Molecule type: mRNA A; Residues: 1-344 (ALL2) A; Cross-references: EMBL:X14355; NID:g3133; PIDN:CAA32536.1; PID:g31334 A; Note: the authors translated the codon ACT for residue 25 as Ala C; Superfamily: Fc gamma receptor I; immunoglobulin homology C; Keywords: alternative splicing; immunoglobulin receptor; transmembrane protein F; 117-170/Domain: immunoglobulin homology C; Maymonds: immunoglobulin homology
	A; Accession: 503019 A; Accession: 503019 A; Residues: 1-344 <all2> A; Residues: 1-344 <all2> A; Residues: 1-344 <all2> A; Cross-references: EMBL:X14355; NID:g31333; PIDN:CAA32536.1; PID:g31334 A; Note: the authors translated the codon ACT for residue 25 as Ala C; Superfamily: Fc gamma receptor I; immunoglobulin homology C; Reywords: alternative splicing; immunoglobulin receptor; transmembrane protein F; 117-170/Domain: immunoglobulin homology <amm></amm></all2></all2></all2>
ma ot	Query Match 100.0%; Score 61; DB 2; Length 344; Best Local Similarity 100.0%; Pred. No. 0.002; Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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P € 6 6 6	FC gamma (196) receptor 1-A (nigh allinity) precursor - numan N.Alternate names: CD64 C;Species: Homo sapiens (man) C;Species: Homo sapiens (man) C;Date: 30-Dec-1991 #sequence_revision 06-Sep-1996 #text_change 21-Jul-2000 C;Date: 30-Dec-1991 #sequence_revision 06-Sep-1996 #text_change 21-Jul-2000 C;Date: 30-Dec-1991 #sequence_revision 06-Sep-1996 #text_change 21-Jul-2000
de te	R;van de Winkel, J.G.J.; Ernst, L.K.; Anderson, C.L.; Chiu, I.M. J. Biol. Chem. 266, 13449-13455, 1991 A. Title: Gene organization of the human high affinity receptor for IgG. FcgammaRI (CD.
us (X	A; Reference number: A39878; MUID:91302383 A; Accession: A39878
ea ·	A; MOJECULE Type: LNA A; Residues: 1-374 < VAN>

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A; Molecule type: mRNA
A; Residues: 1-336 < RE
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Science 260,
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Accession: I57525
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Residues: 1-8 <RE2>
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Reference number: S03018; 1
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Accession: B41357
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esidues: 1-24,'S',26-337,'T',339-374 <AL1>
coss-references: GB:X14356; GB:M21091; NID:g31331;
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78,152,159,163,195,240/Binding site: carbohydrate
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12; Conser
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695-698, 1993
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-336 <RES>
ences: EMBL:X70980; NID:g311748; PIDN:CAA50311.1;
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0.0022;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PIDN: AAD13887.1; PID: g4261587
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PIDN:CAA32537.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        the human high affinity Fc receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               immunoglobulin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 374;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (Asn)
                                                                                                                                                                                    and defective
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  in human mononuclear cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (covalent) #status predict
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PID:g31332
         PID:g311749
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               receptor; transmembrane
                                                                                                                                                                                                                                                P.M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .1; PID:g180279 Kimberly, R.P.
                                                                                                                                                                                       E,C
                                                                                                                                                                                                                                                Wicker,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PID:g31332
                                                                                                                                                                                    receptor
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                                                                                                                                                                                                                                                                                                   R;Alm, R.A.; Ling, L.S.L.; Moir, ; Ives, C.; Gibson, R.; Merberg, Nature 397, 176-180, 1999
                                                                                                                                                                                                                                                                                                                                                                                                                  hydroxyethylthiazole kinase - Helicobacter pylori (strain J99)
C;Species: Helicobacter pylori
A;Variety: strain J99
C;Date: 12-Feb-1999 #sequence_revision 12-Feb-1999 #text_change 21-Jan-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Qy
                                                                        A; Experimental source:
                                                                                                   A; Cross-references:
                                                                                                                              A; Residues:
                                                                                                                                                         A; Molecule type: DNA
                                                                                                                                                                                                                  A; Accession: A71890
                                                                                                                                                                                                                                             A; Title: Genomic sequence comparison of two unrelated isolates A; Reference number: A71800; MUID:99120557
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  F; 127-179/Domain:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    밁
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                                                                                                                                                                                       A;Status: preliminary
                                                                                                                                                                                                                                          A; Reference number: A71800;
                                                                                                                                                                                                                                                                                                                                                                                       C; Accession: A71890
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C;Superfamily: Fc gamma receptor I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Residues:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Molecule type: mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Accession: A43511
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 R;Sears,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C; Accession: A46480;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Residues: 1-404 <OSM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             R;Osman,
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Best Local
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ol. 144, 371-378,
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                                                                                                                        1-259 <ARN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       immunoglobulin receptor; transmembrane protein
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9; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity 9; Conser
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                                                                           :AE001508; GB:AE001439; NID:g4155338; PIDN:AAD06352.1; strain J99
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75.0%;
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D.T.; King, B.L.; D.; Mills, S.D.;

Jiang, Q.; Brown,

Taylor, D.E.;

of the human gastric p

PID:9415

E.D.;

Doig, P.C.;

Smith,

G.F

Score 48; DB Pred. No. 0.49 2; Mismatches

DB 2

2

Length 404;

Indels

0

Gaps

0;

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homology

immunoglobulin

homology

PID:g200753

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C; Superfamily: Fc gamma rec
C; Keywords: immunoglobulin
F; 128-180/Domain: immunoglo
                                                                                                                                                                                                                                                           N;Alternate names: high affinity IgG receptor
C;Species: Mus musculus (house mouse)
C;Date: 18-Jun-1993 #sequence_revision 18-Nov-1994 #text_change 21-Jan-2000
                                                                         A; Note: sequence extracted
                                                                                                                                                  A;Reference number: A46480;
A;Accession: A46480
A; Reference number: A43511;
                   A; Title: Molecular cloning and
                                                                                                                                                                     A; Title: Structure and mapping of the gene encoding A; Reference number: A46480; MUID:92166399
                                                                                                                                                                                                        Immunol. 148, 1570-1575,
                                                                                                                                                                                                                                                                                                                     receptor high affinity - mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  immunoglobulin
                                      8, 1990
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75.0%;
                                                                                                                                                                                                                        McKenzie, I.F.; Hogarth, P.M
                                                                           from
 and expression of MUID:90111035
                                                                                                                                                                                                          1992
                                                        В.;
                                                      NCBI backbone (NCBIN:85205, NCBIN:85208, B.; McKenzie, I.F.C.; Hogarth, P.M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  homology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     I; immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2;
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Pred. No.
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                   the mouse high affinity
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                                                                                                                                                                                     high
                                                                                                                                                                                     affinity Fc
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                     Fc receptor for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                             NCBIN:85212,
                                                                                                                                                                                        gamma
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RIa

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submitted to the EMBL Data Library, A; Reference number: S33979
A; Accession: S33981
                                                                                                                                                                                                                                                                                                                                                                      vif protein - human immunodeficiency virus type 1
C:Species: human immunodeficiency virus type 1, HIV-1
C:Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 20-Sep-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C;Accession: E64625
R;Tomb, J.F.; White, O.; Kerlavage, A.R.; Clayton, R.A.; Sutton, G.G.; Fleischmann, R.D.; Peterson, S.; Loftus, B.; Richardson, D.; Dodson, R.; Khalak, H.G.; Glodek, A.; McKenne son, J.D.; Kelley, J.M.; Cotton, M.D.; Weidman, J.M.; Fujil, C.; Bowman, C.; Watthey, L. Nature 388, 539-547, 1997
Nature 388, 539-547, 1997
A;Authors: Wallin, E.; Hayes, W.S.; Borodovsky, M.; Karpk, P.D.; Smith, H.O.; Fraser, C. A;Title: The complete genome sequence of the gastric pathogen Helicobacter pylori.
A;Reference number: A64520; MUID:97394467
A;Accession: E64625
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Best Local Similarity
Thehes 7; Conserv
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                                                                                                                                                                          A;Cross-references: EMBL:Z11530; C;Superfamily: AIDS vif protein
                                                                                                                                                                                           A; Molecule type: mRNA
A; Residues: 1-192 < CAR>
A; Cross-references: EMB
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                                                                                                                                                                                                                                                                                                                                     R; Carlini,
                                                                                                                                                                                                                                                                                                                                                          C; Accession: S33981
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              hydroxyethylthiazole kinase (EC 2.7.1.50) – Helicobacter pylori (strain 26695)
C;Species: Helicobacter pylori
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Pathway: thiamin biosynthesis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ;Superfamily: hydroxyethylthiazole kinase; hydroxyethylthiazole kinase homology;Reywords: phosphotransferase; thiamin biosynthesis;16-203/Domain: hydroxyethylthiazole kinase homology <HTK>
                                                                                                                                                                                                                                                       Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Description: catalyzes the phosphorylation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity Matches 7; Conserv
                                                                                                Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local
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126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    134 GSLVGISCESK 144
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                                       1 GNLVTLSCE 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 GNLVTLSCETK 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GNLVTLSCETK 11
GNVVRLSCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          preliminary; nucleic acid sequence not shown; translation not shown
                                                                                                Similarity
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                                                                          Conservative
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134
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                                                                                              Score
Pred.
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                                                                              Mismatches
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No.
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15;
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13;
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14;
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                                                                                                                 Length 192
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hypothetical protein DKFZp C. Species: Homo sapiens (m C. Species: 11-Jun-1999 #seque C; Accession: T08678 R. Wambutt, R.; Heubner, D. submitted to the Protein S. A.; Reference number: Z16469 A; Accession: T08678
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C;Species: Drosophila virilis
C;Date: 30-Jun-1992 #sequence_revision 30-Jun-1992 #text_change 21-Feb-1997
C;Accession: A41544
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Gene: FlyBase:sina
A;Cross-references: FlyBase:FBgn0003410
C;Keywords: DNA binding; nucleus; zinc finger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Title: seven in absentia, a gene required for specification of R7 cell fate in the A;Reference number: A36195; MUID:91029488
A;Accession: A36195
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C; Species: Drosophila melanogaster
C; Date: 14-Dec-1990 #sequence_revision 13-Jan-1993 #text_change 24-Sep-1998
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C;Keywords: DNA binding; zinc finger
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C;Genetics:
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A;Molecule type: DNA
A;Residues: 1-314 <NEU>
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A; Title: Evolution of gene position: chro
A; Reference number: A41544; MUID:92052239
A; Molecule type: mRNA
A; Residues: 1-584 <WAM>
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A; Residues: 1-314 <CAR>
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                                                                                                                                                                                  protein DKFZp564I1922.1 - human (fragment)
                                                                                                  Heubner, D.; Mewes,
                                                                                                                                         1999 #sequence_revision 11-Jun-1999 #text_change 23-Jul-1999
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Pred. No. 24;
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se, May 1999
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RESULT 13
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A;Experimental source: fetal brain
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A;Accession: A32578
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C;Date: 21-May-1990 #sequence_revision 21-May-1990 #text_change 30-May-1997
C;Accession: A32578
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A;Title: Diversity of T cell receptor alpha and beta chain genes expressed by human T ce A;Reference number: S57869; MUID:92192091
                                                                                                                                                                                                                                                                                                                                                                                                                                     C; Species: Homo sapiens
C; Date: 27-Oct-1995 #sec
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C;Superfamily: immunoglobulin V
C;Keywords: T-cell receptor
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A; Residues: 1-113 <KIM>
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Residues: 1-137 <GIE>
Superfamily: immunoglobulin
Keywords: T-cell receptor
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J. Immunol. 17, 375-383, 1987
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   receptor beta chain precursor ies: Homo sapiens (man)
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                                                                                                                   hypothetical protein slr1243 - Synechocystis sp. (strain PCC 6803)
C.Species: Synechocystis sp.
A; Variety: PCC 6803
C; Date: 25-Apr-1997 #sequence_revision 25-Apr-1997 #text_change 20-Jun-2000
C; Accession: S74861
R; Kaneko, T.; Sato, S.; Kotani, H.; Tanaka, A.; Asamizu, E.; Nakamura, Y.; A.
O, K.; Okumura, S.; Shimpo, S.; Takeuchi, C.; Wada, T.; Watanabe, A.; Yamada
DNA Res. 3, 109-136, 1996
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S74861
      A;Reference number: $74322;
A;Accession: $74861
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-1064 <KAN>
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A;Title: Sequence analysis of the genome of
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364

MUID:97061201

the unicellular cyanobacterium

Synechocys

Yamada,

Yas

2 NLVTLSCETKL 12

Conservative

54.5%;

Score 37; DB Pred. No. 43; 2; Mismatches

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Length 382

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C; Superfamily: immunogiobile C; Keywords: T-cell receptor
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R;Jores, R.; Meo, T.
submitted to the EMBL Data Library,
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    A; Gene: MTH141
C; Superfamily:
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A;Status: preliminary
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                                                                    A;Cross-references: GB:AE000903; GB:AE000666; NID:g2622514; A;Experimental source: strain Delta H
                                                                                                                                                                                                                                                                                                                                                                                    probable cell division control protein 6^{-\epsilon} Methanobacterium thermoautotrophicum (stra C; Species: Methanobacterium thermoautotrophicum
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                                                                                                                                                                                                     J. Bacteriol. 179, 7135-7155, 1997
A;Title: Complete genome sequence of Methanobacterium thermoautotrophicum Delta H:
A;Reference number: A69000; MUID:98037514
                                                                                                                                                                                                                                                                                                                       R;Smith, D.R.;
                                                                                                                                                                                                                                                                                                                                            C; Accession:
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                                                                                                                                        A; Molecule type: DNA
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Archaeoglobus fulgidus probable cell
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79, 7135-7155, 1997
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A;Cross-references: EMBL:D90909; GB:AB001339; NID:g1652844; PIDN:BAA17822.1; PID:g165290 A;Note: the nucleotide sequence was submitted to the EMBL Data Library, June 1996 C;Superfamily: Synechocystis hypothetical protein s1r1243
Query Match 60.7%; Score 37; DB 2; Length 1064; Best Local Similarity 45.5%; Pred. No. 1.1e+02; Matches 5; Conservative 4; Mismatches 2; Indels
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Q99795	014953	P43631	Q9x5z9	031213	Q10343	P54356	Q62230	090703	015394	P27462	P04217
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ALIGNMENTS

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• • • • • • • • • • • • • • • • • • • •	<pre>entities requires a license agreement (See http://www.isb-sib.ch/announce/ or send an email to license@isb-sib.ch).</pre>	ified and this statement is not removed. Usage by and for commerc	by non-profit institutions as long as its content is in no	the European Bioinformatics Institute of Bioinformatics and the EMBL outstation -	LSS-PROT entry is copyright. It is produced through		WWW-"http://www.ncbi.nlm.nih.gov/prow/cd/cd64.htm".	-1- DATABASE: NAME-PROW; NOTE-CD guide CD64 entry;	-!- SIMILARITY: CONTAINS 3 IMMUNOGLOBULIN-LIKE C2-TYPE DOMAINS.	-!- TISSUE SPECIFICITY: MONOCYTE/MACROPHAGE-SPECICIC.	ED BY ALTERNATIVE SPLICING.	PRODUCTS: 2 ISC	-!- SUBCELLULAR LOCATION: TYPE I MEMBRANE PROTEIN.	AFFINITY RECEPTOR.	- 1- FUNCTION BINDS TO THE FC REGION OF IMMUNOSIOBULINS GAMMA HIGH	complementary DNAS."; Science 243:378-381/1989)	"Isolation and expression of functional high-affinity Fc receptor			SEQUENCE FROM N.A.		Nucleic Acids Res. 16:11824-11824(1988).		"Nucleotide sequence of three cDNAs for the human high affinity Fc			SEQUENCE FROM N.A.	(1)		Mammalia: Eutheria: Primates: Catarrhini: Hominidae: Homo.	Chordata: Craniata: Vortobrata:	FOSTA ON FOST ON ECST ON ISERT.	RI) (FCRI) (IGG FC RECEPTOR I) (CD64).	ITY IMMUNOGLOBULIN G	te)	(Rel. 12,	01-OCT-1989 (Rel. 12, Created)	2315;	FCG1_HUMAN STANDARD; PRT; 374 AA.	FCG1_HUMAN	LT 1

EMBL; X14356; CAA32537.1; -. EMBL; X14355; CAA32536.1; -.

PIR; A41357; A41357 PIR; B41357; B41357 HSSP; P12319; 1ALT. PIR; S03018; S03018 PIR; S03019; S03019 PIR; A41357; A41357

HSSP; P12319 MIM; 146760;

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J. Immunol. 148:1570-1575(1992).

-I- FUNCTION: BINDS TO THE FC REGION OF IMMUNOGLOI AFFINITY RECEPTOR.

-I- SUBCELLULAR LOCATION: TYPE I MEMBRANE PROTEIN.

-I- TISSUE SPECIFICITY: MACROPHAGE-SPECICIC.

-I- SIMILARITY: CONTAINS 3 IMMUNOGIORITY TABLETTER.
                                                                                                                                                                                                                                                                                                                                                                                                                             FCG1_MOUSE STANDARD; PRT; 404 AA P26151; PRT; 404 AA 01-MAY-1992 (Rel. 22, Created) 01-MAY-1992 (Rel. 22, Last sequence update) 15-JUL-1999 (Rel. 38, Last annotation update) HIGH AFFINITY IMMUNOGLOBULIN GAMMA FC RECEPTION
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
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Osman N., Kozak C.A., McKenzie I.F.,
                                                                                                      "Structure and mapping of the gene gamma RI and chromosomal location of
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MEDLINE-99120557; PubMed=9923682;

Alim R.A., Ling L.-S.L., Moir D.T., King B.L., Brown E.D., Doi

Smith D.R., Noonan B., Guild B.C., deJonge B.L., Carmel G.,

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01-OCT-2000 (Rel. 40, Last annotation update)
HYDROXYETHYLTHIAZOLE KINASE (EC 2.7.1.50) (4-
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PIR; A43511; A43511.
PIR; A46480; A46480.
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InterPro; IPR003006; -
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gastric pathogen Helicobacter pylori.";
nature 397:176-180(1999).
-i- CATALTIC ACTIVITY: ATP + 4-METHYL-5-(2-HYDROXYETHYL)-THIAZOLE
ADP + 4-METHYL-5-(2-PHOSPHOETHYL)-THIAZOLE.
-i- COFACTOR: MAGNESIUM (BY SIMILARITY).

"Genomic sequence comparison

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HYDROXYETHYLTHIAZOLE KINASE (EC 2.7.1.50) (4-METHYL-5-BETA-
HYDROXYETHYLTHIAZOLE KINASE) (THZ KINASE) (TH KINASE).
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MEDLINE=97394467; Pu
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                                                                                                                                                                                            -1- COFACTOR: MAGNESIUM (BY SIMILARITY).
                                                                                                                                                                                                                                                                 Nature 388:539-547(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=210;
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SIMILARITY: BELONGS TO THE THZ KINASE FAMILY.
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BELONGS TO THE THZ KINASE FAMILY.
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                                     RA Beson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S.,
RA Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P.,
RA Burtis K.C., Busam D.A., Butler H., Cadieu E., Center A., Chandra I.,
RA Burtis K.C., Busam D.A., Butler H., Cadieu E., Center A., Chandra I.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Dietz S.M.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Dietz S.M.,
RA Dodson K., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P.,
RA Durbin K.J., Evangelista C.C., Ferraz C., Ferriera S., Fleischmann W.,
RA Fosler C., Gabrielian A.E., Garg N.S., Gelbart W.M., Glasser K.,
RA Glodek A., Gong F., Gorrell J.H., Gu Z., Guan P., Harris M.,
RA Harris N.L., Harvey D., Heiman T.J., Hernandez J.R., Houck J.,
RA Hostin D., Houston K.A., Howland T.J., Wei M.-H., Ibegwam C.,
RA Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z.,
RA Lasko P., Lei Y., Levitsky A.A., Li J., Liang Y., Lin X.,
RA Morkulton K. Maltei B., McIntosh T.C., McLeod M.P., McPherson D.,
RA Morkulton K. Maltei B., McIntosh T.C., McLeod M.P., McPherson D.,
RA Markulton K. Milehina N. W. McPherson D., McPherson D.,
RA Markulton K. Milehina N. W. McPherson D., McPherson D.,
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                                                                                                                                                                                                                                                                                                                                                                                                            Abril J.F., Agbayani A., An H.-J., Andrews-Pfannkoch C., Baldwin D. Ballew R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M., Beesson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N., Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X., Brandon R.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfeiffer B.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-MAY-1991 (Rel. 18; Created)
01-NOY-1995 (Rel. 32; Last sequence update)
01-CCT-2000 (Rel. 40; Last annotation updat
DEVELOPMENTAL PROTEIN SEVEN IN ABSENTIA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE=20196006; PubMed=10731132;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRAIN-BERKELEY;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE=91029488; PubMed=2146028; Carthew R.W., Rubin G.M.;
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Neoptera; Endopterygo
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MAGNESIUM (BY SIMILARITY
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                             Morris J
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Spier E., Spr
Svirskas R.,
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01-DEC-1992 (Rel. 24, Last sequence update)
01-OCT-2000 (Rel. 40, Last annotation updat
DEVELOPMENTAL PROTEIN SEVEN IN ABSENTIA.
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SEQUENCE
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"The genome sequence of Drosophila melanogaster.";
Proc. Natl. Acad. Sci. U.S.A. 88:10203-10207(1991).
-!- FUNCTION: REQUIRED FOR SPECIFICATION OF R7 PHOTORECEPTOR CELL FATE IN THE DROSOPHILA EYE. POSSIBLY ACTS BY REGULATING GENE
                                                                                              MEDLINE=92052239; PubMed=1946441;
Neufeld T.P., Carthew R.W., Rubin G.M.;
"Evolution of gene position: chromosomal arrangement and
comparison of the Drosophila melanogaster and Drosophila
                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Arthropoda; Tracheata; Pterygota; Neoptera; Endopterygota; Diptera
Ephydroidea; Drosophilidae; Drosophila.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   modified and this statement is not removed entities requires a license agreement (See or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                     Drosophila virilis 
Eukaryota; Metazoa;
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the European Bioinformatics Institute
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EM
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                                                                                                                                                                                              SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Reinert K.,
                                                                                                                                                                                                                                             NCBI_TaxID=7244;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Palazzolo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     90
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FUNCTION: REQUIRED FOR SPECIFICATION OF R7 PHOTORECEPTOR CELL FATE IN THE DROSOPHILA EYE. POSSIBLY ACTS BY REGULATING GENE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EXPRESSION.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GHLVCVSCRSKL 101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AE003526; AAF49403.1;
                                                                       genes."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FBgn0003410; sina
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             non-profit institutions as long
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             , Siden-Kiamos I., Simpson M., Skupski M.P., Smit
Spradling A.C., Stapleton M., Strong R., Sun E.
R., Tector C., Turner R!, Venter E., Wang A.H., V
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   314 AA;
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, Pittman G.S., 'Pan S., Pollard J., Puri V.
Remington K., Saunders R.D.C., Scheler F.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein;
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                                                                                                                                                                                                                                                                                                                                           (Fruit fly).
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33707
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58.3%;
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, 'Pan S., Pollard J., Puri V.,
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TO DICTYOSTELLIUM DG17 (AA 25-67).
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                                                                                                                                                                                                                                                                                                                                                                                          ABSENTIA.
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                                                                                                                                                                                                                                                                                            Diptera;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Reese M.G.,
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- outstation
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RESULT 7
C2TA_MOUSE
      ALD DEPTH OF THE PROPERTY OF T
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Best Local
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Cancer Immunol. Immunother. 42:99-107(1996).

Cincer Immunol. Immunother. 42:99-107(1996).

Cincer Immunol. Immunother. 42:99-107(1996).

CIPROMOTER: ACTIVATION IS VIA THE PROXIMAL ACTIVITY OF THE HLA CLIP CONCERNING OF IN VITRO TRANSLATED CIITA WAS DETECTED. MAY ACT IN COACTIVATOR-LIKE FASHION THROUGH PROTEIN INTERACTIONS CONCERNING THE COACTIVATOR COA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "Mouse class II transactivator: cDNA sequence and amino comparison with the human class II transactivator."; Immunogenetics 45:220-222(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C2TA_MOUSE STANDARD; PRT; 1078 AA. P79621; Q31115; C1eated)
15-DEC-1998 (Rel. 37, Created)
15-DEC-1998 (Rel. 37, Last sequence update)
01-OCT-2000 (Rel. 40, Last annotation update)
MHC CLASS II TRANSACTIVATOR CLITA.
MHC2TA OR CLITA OR C2TA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE=97152412; PubMed=8995190; Sims T.N., Elliott J.F., Ramassar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mus musculus (Mouse).
Eukaryota; Metazoa; (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRAILBEALB/C; TISSUE=Thymus;
MEDLINE=9618886; PubMed-862527;
Panelli M.C., Wang E., Shen S., S
Hersh E.M., Stopeck A., Gangavall
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EMBL; M77282; AAA28899.1; -. PIR; A41544; A41544.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         the European Bioinformatics Institute.
                                                                                                                                                                                                                                                                                                                                                                                  poorly responsive to IFNgamma
                                                                                                                                                                                                                                                                                                                                                                                                                                         Akporiaye E.T.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   between
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE OF 878-1020 FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NCBI_TaxID=10090;
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                                                                                                                                                                                                                                                                                                                                    mmunogenicity is accompanied by induction
                                                                                                                                                                                                                                                                                                                                                                                                                 Interferon gamma (IFNgamma) gene transfer of an EMT6 tumor that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         90 GHLVCVSCRSKL 101
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CONTACTING FACTORS BINDING TO THE PROXIMAL MHC CLASS TO ELEMENTS OF THE TRANSCRIPTION MACHINERY, OR BOTH-IT MAY ACTIVATE HLA CLASS II TRANSCRIPTION BY MODITY THAT BIND TO THE MHC CLASS II PROMOTER (BY SIMILARITY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SUBCELLULAR LOCATION: NUCLEAR.
TISSUE SPECIFICITY: IN MANY OMMATIDIAL PRECURSOR CELLS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity 7; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       non-profit institutions as long and this statement is not removed.
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k A., Gangavalli
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ramassar V.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WW.
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Pred. No.
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TO DICTYOSTELLIUM DG17 (AA 25-67).
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lli R., Barber
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J., Jolly D.,
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                                                                                                                                                                                                                                                                                                                        Nagase T., Seki N., Ishikawa K.-I., Ohira M., Kawarabayasi Y., Ohara O., Tanaka A., Kotani H., Miyajima N., Nomura N.; "Prediction of the coding sequence of unidentified human genes. The coding sequences of 80 new genes (KIAA0201-KIAA0280) deduced analysis of cDNA clones from cell line KG-1 and brain.";
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                                                                                                                                                                                       or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                the European Bioinformatics Institute.
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MEDLINE-97191544; PubMed-9039502;
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Mammalia; Eutheria;
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Transcription regulation;
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15.JUL-1998 (Rel. 36, Created)
15.JUL-1998 (Rel. 36, Last sequence update)
11-JUL-1998 (Rel. 36, Last sequence update)
01-OCT-2000 (Rel. 40, Last annotation update)
INTERLEUKIN-18 PRECURSOR (IL-18) (INTERFERON-CAMMA INDUCING FACTOR)
(IFN-GAMMA-INDUCING FACTOR) (INTERLEUKIN-1 GAMMA) (IL-1 GAMMA).
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T-CELL SURFACE GLYCOPROTEIN CD4 PRECURSOR (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE
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between the Swiss Institute of Bioinformatics and the EMBL
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FUNCTION: AUGMENTS NATURAL KILLER CELL ACTIVITY IN SPLEEN CELLS AND STIMULATES INTERFERON GAMMA PRODUCTION IN THELPER TYPE I
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139
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                                                                                                                                                                                                                                                                                                                                                    Score 36;
Pred. No.
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MS -> IP (IN REF. 2).

I -> M (IN REF. 2).

; E2089AD6F1798450 CRC64;
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RESULT 11
VPL_HAEIN
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Best Local
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                              01-NOV-1995 (Rel. 3
01-NOV-1995 (Rel. 3
01-OCT-2000 (Rel. 4
MU-LIKE PROPHAGE FI
                                          01-NOV-1995
01-NOV-1995
01-OCT-2000
                                                                        VPL_HAEIN P44233;
                                                                                                                                                                                                                                                                           DISULFID DISULFID
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SIGNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Submitted (AUG-1996) to the EMBL/GenBank/DDBJ databases.
-I- FUNCTION: ACCESSORY PROTEIN FOR MHC CLASS-II ANTIGEN/T-CELL RECEPTOR INTERACTION. MAY REGULATE T-CELL ACTIVATION.
-I- SUBUNIT: ASSOCIATES WITH P56-LCK (BY SIMILARITY).
-I- SUBCELLULAR LOCATION: TYPE I MEMBRAME PROTEIN.
-I- SUBCELLULAR LOCATIONS TO THE IMMUNOGLOBULIN SUPERFAMILY. CONTAINS ONE V-LIKE AND 3 C2-LIKE DOMAINS.
 Haemophilus
Bacteria; Pr
                                                                                                                                                                                                                                                                                                                                                                                                                                Pfam; Processing CD41C.
                                                                                                                                                                                                                                   SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  entities requires a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         use by non-profit institutions as long modified and this statement is not removed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Ve
Mammalia; Eutheria; Primates; Platyrrhini;
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                                                                                                                                                                                                                                                       LIPID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             the European Bioinformatics Institute.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tatsumi M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NCBI_TaxID=9521;
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                                                                                                                                      GEVVELPCETSL
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                                                                                                                                                           GNLVTLSCETKL
                                                                                                                                                                                Similarity 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                           Lipoprotein;
  Proteobacteria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IPR000973; -.
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            influenzae
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                                                                                    STANDARD;
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                              32, Created)
32, Last sequence update)
40, Last annotation update)
FLUMU TAIL SHEATH PROTEIN.
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Transmembrane;
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gamma
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BY SIMILAR
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PALMITATE
PALMITATE
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                                                                                                                                                                                                                                                                                                                                                                        POTENTIAL.
CYTOPLASMIC (POTENTIAL)
                                                                                                                                                                                                                                                                                                                                                                                           EXTRACELLULAR
                                                                                    PRT;
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ALMITATE (BY SIMILARITY).
57,EED6344005A015 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                SIMILARITY.
subdivision; Pasteurellaceae;
                                                                                                                                                                                                                                                                 SIMILARITY
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D (GLCNAC. . .)
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ni; Cebidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   as
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                                                                                                                                                                                                  Length 457;
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Cebinae; Saimiri.
                                                                                                                                                                                                                                                                                              (POTENTIAL).
                                                                                                                                                                                                                                                                                                                                                                                                                                    T-cell; MHC;
                                                                                                                                                                                                                                                                                                                   (POTENTIAL).
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D SAHZ_INVILL

C P50245; Q27587; Q9VEQ4;

C P50245; Q27587; Q9VEQ4;

OT 01-CCT-1996 (Rel. 34, Created)

OT 01-CCT-1996 (Rel. 34, Last sequence update)

OT 01-CCT-2000 (Rel. 40, Last annotation update)

PUTATIVE ADENOSYLHOMOCYSTEINASE (EC 3.3.1.1) (S-ADENOSYL-L-

HOMOCYSTEINE HYDROLASE) (ADDHCYASE).

BE HOMOCYSTEINE HYDROLASE) (ADDHCYASE).

GN AHCY89E OR PH200 OR CG8956.

OS Drosophila melanogaster (Fruit fly).

CE Bukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;

CC Eukaryota; Metazoa; Arthropoda; Diptera; Brachycera; Musco

CC Pterygota; Neoptera; Endopterygota; Diptera; Brachycera; Musco
     ACC CRAFT RACE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              В
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Best Local s
Matches 6
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MEDLINE=95396803; PubMed=7667301;
Martin C.H., Mayeda C.A., Davis C
Mathog D.R., Celniker S.E., Lewis
                                                    SEQUENCE FROM N.A. STRAIN=CANTON-S;
                                                                                                     trans-regulatory homeoprotein.";
EMBO J. 7:3223-3231(1988).
                                                                                                                               MEDLINE-89030618; PubMed-2903049;
Delorenzi M., Ali N., Saari G., Henry C., Wilcox M.,
Delorenzi M., Ali N., Saari G., Henry C., Wilcox M.,
"Evidence that the Abdominal-B r element function is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       use by non-profit institutions as long modified and this statement is not removed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Science 269:496-512(1995).
-!- FUNCTION: MAJOR COMPONENT OF
-!- SIMILARITY: STRONG, TO PHAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=95350630; PubMed=7542800;
                                                                                                                                                                                            TISSUE=Embryo;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          SAH2_DROME
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          entities requires a
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NCBI_TaxID=727;
                                                                                                                                                                                                              SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "Whole-genome random sequencing and assembly of Haemophilus influenzae \operatorname{Rd}.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        tities requires a license agreement (S send an email to license@isb-sib.ch).
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by non-profit institutions as long as its content
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity 54.96; Conservative
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487 AA; 53143
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                Ericsson C.L.,
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Lewis

Е.В.,

Palazzolo M.J.;

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RA Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S.,
RA Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P.,
Burtis K.C., Busam D.A., Butler H., Cadieu E., Center A., Chandra I.,
RA Burtis K.C., Busam D.A., Butler H., Cadieu E., Center A., Chandra I.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Dodson K., Doup L.E., Downes M., Dugan-Rocha S., Dlukov B.C., Dunn P.,
RA Durbin K.J., Evangelista C.C., Ferraz C., Ferriera S., Fleischmann W.,
RA Posler C., Gabrielian A.E., Garg N.S., Gelbart W.M., Glasser K.,
RA Flock A., Gong F., Gorrell J.H., Gu Z., Guan P., Harris M.,
RA Glodek A., Gong F., Gorrell J.H., Gu Z., Guan P., Harris M.,
RA Harris N.L., Harvey D., Heiman T.J., Hernandez J.R., Houck J.,
RA Harris N.L., Harvey D., Heiman T.J., Wei M.-H., Ibeywam C.,
RA Hostin D., Houston K.A., Howland T.J., Wei M.-H., Ibeywam C.,
RA Jalai M., Kalush F., Karpen G.H., Ke Z., Keulp D., Lai Z.,
RA Lasko P., Lei Y., Levitsky A.A., Li J., Li Z., Liang Y., Lin X.,
RA Lasko P., Lei Y., Levitsky A.A., Li J., McDherson D.,
Marteli B., McIntosh T.C., McLeod M.P., McPherson D.,
Martin D., McIntosh T.C., McLeod M.P., McPherson D.,
McPherson D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Merkulov G., Milshina N.V., Mobarry C., Morris J., Moshrefi A., A Mount S.M., Moy M., Murphy B., Murphy L.M., Muzny D.M., Nelson D.L., Wa Nelson D.R., Pacleb J.M., Wa Palazzolo M., Pittman G.S., Pan S., Pollard J., Puri V., Reese M.G., Remington K., Sannders R.D.C., Scheeler F., Shen H., Shue B.C., Siden-Klamos I., Simpson M., Skupski M.P., Smith T., Wa Spier E., Spradling A.C., Stapleton M., Strong R., Sun E., Wang A.C., Tector C., Turner R., Venter E., Wang A.H., Wang X., Wang Z.-Y., Wassarman D.A., Weinstock G.M., Weissenbach J., Walliams S.M., Woodage T., Worley K.C., Wu D., Yang S., Yao Q.A., Ye J., Yeh R.-F., Zaveri J.S., Zhan M., Zhang G., Zhao Q., Zheng L., Zheng X.H., Zhong F.N., Zhong M., Zhou X., Zhu X., Smith H.O. A Zheng R.A., Myers E.W., Rubin G.M., Venter J.C., The Genome sequence of Drosophila melanogaster."; Science 287:2185-2195(2000).
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                                                                                                                                                                                                                                                                            EMBL; U31961; AAA84400.1; -.
EMBL; AE003715; AAF55367.1; -.
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                                                                                                                           PROSITE; PS00738; ADOHCYASE_1;
PROSITE; PS00739; ADOHCYASE_2;
                                                                                                                                                                                                                                                FlyBase; FBgn0015011; Ahcy89E.
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                                                               NP_BIND
                                                                                              Hydrolase;
                                                                                                                                                                                                                          InterPro;
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SIMILARITY: BELONGS TO THE ADENOSYLHOMOCYSTEINASE FAMILY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PATHWAY: ACTIVATED METHYL CYCLE..
TISSUE SPECIFICITY: EXPRESSED IN EXTENDI
SOMATIC MESODERM, YOLK CELLS AND MIDGUT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CATALYTIC ACTIVITY: S-ADENOSYL-L-HOMOCYSTEINE
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NAD;
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                                                                                              metabolism
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   NAD (POTENTIAL).
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LATER EMBRYONIC
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Zhu X., Smith H.O.
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070472;
01-0CT-2000
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                                                                                                         P13944: Q04509;
01-JAN-1990 (Rel. 13, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
01-OCT-2000 (Rel. 40, Last annotation update)
COLLAGEN ALPHA 1(XII) CHAIN PRECURSOR (FIBROCHIMERIN).
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                                                                                                                                                                                                                                                                                                                                                       EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                     between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
SEQUENCE FROM N.A. STRAIN-WHITE LEGHORN;
                                                                       Gallus gallus (Chicken).
Eukaryota; Metazoa; Chordata;
                                                                                                                                                                                   CHICK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Wilkinson R., Fitter S., Tscharke D., Simmons A.;
Submitted (APR-1998) to the EMBL/GenBank/DDBJ databases.
-!- FUNCTION: MAY PLAY A ROLE IN THE IMMUNE RESPONSE TO VIRAL
                                                           Archosauria;
                                                                                                 COL12A1
                                                                                                                                                                       CA1C_CHICK
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                                                           Craniata; Vert; Galliformes;
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22;
                                                                         Vertebrata; Euteleostomi;
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                                                             Phasianidae;
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                                                              Phasianinae;
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Yamaqata M., Yamada K.M., Ishinana Yamaqata M., Yamada K.M., Ishinana K.;;
Nishida Y., Obara M., Kimata K.;;
"The complete primary structure of type XII collagen shows a chimeric molecule with reiterated fibronectin type III motifs, von Willebrand factor A motifs, a domain homologous to a noncollagenous region of factor A motifs, a domain homologous to a noncollagenous region of type IX collagen, and short collagenous domains with an Arg-Gly-Asp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "Large and small splice variants of collagen XII: differential expression and ligand binding."; [
J. Cell Baol. 130:1005-1014(1995).

-1- FUNCTION: TYPE XII COLLAGEN INTERACTS WITH TYPE I COLLAGEN-
CONTAINING FIBRILS, THE COLL DOMAIN COULD BE ASSOCIATED WITH THE
SURFACE OF THE FIBRILS, AND THE COL2 AND NC3 DOMAINS MAY BE
LOCALIZED IN THE PERIFIERILLAR MATRIX.

-1- SUBUNIT: TRIMER OF IDENTICAL CHAINS EACH CONTAINING 190 KDA OF
NORMEDTY: TRIMER OF IDENTICAL CHAINS EACH CONTAINING 190 KDA OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Trueb J., Trueb B.;
"The two splice variants of collagen XII share a Biochim. Biophys. Acta 1171:97-98(1992).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "Type XII collagen: distinct extracellular matrix component discovered by cDNA cloning."; Proc. Natl. Acad. Sci. U.S.A. 84:6040-6044(1987).
                            use by non-profit institutions as long as modified and this statement is not removed. U
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                                                                                          the European
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MEDLINE=93042014; PubMed=1420368;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gordon M.K., Gerecke D.R., Dublet B., van "Type XII collagen. A large multidomain mc homology to type IX collagen.";
J. Biol. Chem. 264:19772-19778(1989).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Koch M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ALTERNATIVE SPLICING.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE OF 1-1283
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                                                                                                                                                                                                                                                                                                                                                              EACH END.
PTM: PROLINE:
UNIT (G-X-Y)
                                                                                                                                                                                           PYM: GLYCOSYLATED, CHONDROITIN-SULFATED (BY SIMILARITY).
SIMILARITY: CONTAINS 4 VWFA DOMAINS.
SIMILARITY: CONTAINS 18 FIBRONECTIN TYPE III-LIKE DOMAINS.
SIMILARITY: BELONGS TO THE FAMILY OF FIBRIL-ASSOCIATED COLLAGENS
WITH INTERRUPTED HELICES (FACIT).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SUBUNIT: TRIMER OF IDENTICAL CHAINS EACH CONTAINING 190 KDA OF NONTRIPLE-HELICAL SEQUENCES.

ALTERNATIVE PRODUCTS: 2 ISOFORMS; A LONG FORM (SHOWN HERE) AND A SHORT FORM; ARE PRODUCED BY-ALTERNATIVE SPLICING. THE FINAL TISSUE FORM OF COLLAGEN XII MAY CONTAIN HOMOTRIMERS OF EITHER THE LONGER OR THE SHORTER ISOFORM OR ANY COMBINATION OF LONG AND SHORT ISOFORM CHAINS. ONLY. THE LONG VARIANT IS A PROTEOGLYCAN. THE LARGE ISOFORM HAS MORE RESTRICTED EXPRESSION IN EMBRYONIC TISSUE THAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                PTM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LIGAMENTS, PERICHONDRIUM, AND PERIOSTEUM, ALL DENSE CONNECT. TISSUES CONTAINING TYPE I COLLAGEN.

DOMAIN: THIS SEQUENCE DEFINES FIVE DISTINCT DOMAINS, TWO TR. HELICAL DOMAINS (COL1 AND COL2) AND THREE NONTRIPLE-HELICAL DOMAINS (COL1 AND COL2).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TISSUE SPECIFICITY: TYPE XII COLLAGEN IS PRESENT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              THE SMALL
                                                                                                                SWISS-PROT entry is copyright. It is produced through een the Swiss Institute of Bioinformatics and the EN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        95370352; PubMed=7642694; Bohrmann B., Matthison M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                         THE TRIPLE-HELICAL
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                                                                                 Swiss Institute of Bioinfo
Bioinformatics Institute.
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                                                                                                                                                                                                                                                                                                                                                                  ARE HYDROXYLATED IN SOME OR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PubMed=2584192;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            TAIL
                                                                                                                                                                                                                                                                                                                                                                                                                                                         IS STABILIZED
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MEDLINE=88166903; PubMed=3127222;
Brueggemann M., Delmastro-Galfre P., Waldmann H., Calabi F.;
Brueggemann M. Delmastro-Galfre P., Waldmann H., Calabi F.;

"Sequence of a rat immunoglobulin gamma 2c heavy chain constant region cDNA: extensive homology to mouse gamma 3.";

Eur. J. Immunol. 18:317-319(1988).
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
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Copyright (c) 1993 - 2000 Compugen Ltd.
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                                                                                                                                                                                                                                                                                                                               MEDLINE=93055454; PubMed=1430234; Porges A.J., Redecha P.B., Doebele R., Kimberly R.P.;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                            EMBL; L03418; AAA36049.1;
HSSP; P12319; FALT.
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Pred. No. 0.0003;
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O58394 pyrococcus
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O911j2 streptomyce
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Best Local :
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"Molecular cloning and identification of full-length cDNA encoding high affinity Fc receptor for bovine IgG (Fc gamma RI)."; Vet. Immunol. Immunopathol. 75:151-159(2000).
EMBL; AF162866; AAF80477.1; -.
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J. Exp. .
791, M9155
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NCBI_TaxID=9913;
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Mammalia; Eutheria; Cetartiodactyla;
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Q38131;
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PROSITE; I
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01-NOV-1996 (TrEMBLrel.
01-NOV-1998 (TrEMBLrel.
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Halocynthia roretzi (Sea squirt).
Eukaryota; Metazoa; Chordata; Urochordata;
Stolidobranchia; Pyuridae; Halocynthia.
      lactococcal bacteriophage rlt.";
Mol. Microbiol. 19:1343-1355(1996)
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                                         MEDLINE=96332669;
van Sinderen D.,
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Nauta A., van Sinderen
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                        Venema G., Nauta A.;
"Sequence analysis and molecular characterization
                                                                                      isolated
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                                                             from Lactococcus lactis bacteriophage rlt."; robiol. 19:1331-1341(1996).
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                                    Park J.-H., Jensen B.C., Pars
"A novel nucleolar G-protein
Submitted (SEP-1999) to the E
EMBL; AF189284; AAF01061.1; -
INTEMPRO; IPR000765; -
                                                                                                                                                                                                                                                                           NOG1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human immunodeficiency virus type 1.
Viruses; Retroid viruses; Retroviridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZNE
                                                                                                                                                STRAIN-ISTAR1;
                                                                                                                                                                                                      Eukaryota; Euglenozoa;
NCBI_TaxID=5691;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "HIV-1 env sequence variation in brain related neurologic disease.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A. MEDLINE-92092169; PubMed-1684385;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ENVELOPE
                                                                                                                                                                                                                                                                                              NUCLEOLAR G-PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Envelope protein.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=11676;
                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                   rypanosoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PAM; PF00516; GP120; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GNLVTLSCETK 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GNLVTLSCETK 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        M75003; AAB08915.1;
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8; Conserv
                         PR00326;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           r. Immune Defic. Syndr. 4:1082-1092(1991).
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                                                                                                                                                                                                                                                                                                             (TrEMBLrel. 13, TrEMBLrel. 13, (TrEMBLrel. 14,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               64 AA;
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  ΑA;
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GTP1OBG.
A; 74797 MW;
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72
                                                                                                                                                                                                                               Kinetoplastida;
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                                                                                 conserved in eukaryotes.";
EMBL/GenBank/DDBJ databases
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Last annotation updat
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Pred.
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Pred.
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  3244326657BDA3E6 CRC64;
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brain tissue of patients with AIDS-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              No.;
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                                                                                                                                                                                                                               Trypanosomatidae;
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Query Match
Best Local S
Matches 6
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SEQUENCE FROM N.A.
SEQUENCE FROM N.A.
Amano H., Kato K., Miyamura T.
Amano H., Kato K., Miyamura T.
Sequence analysis of Yaba vir
Submitted (MAR-1999) to the EV
EMBL; AB025319; BAA88969.1;
EMBL; AB025319; BAA88969.1;
                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                  Query Match
Best Local S
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01-OCT-2000
01-OCT-2000
01-OCT-2000
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01-MAY-2000
01-MAY-2000
01-MAY-2000
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                                                                                                                                      Viruses; dsDNA viruses,
                                                                                                                                               Yaba monkey tumor virus
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                                                                                                                                                                                                                                                                                                                                                                    EMBL;
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                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A!
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRAIN-COLUMBIA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Arabidopsis thaliana (Mouse-ear cress).
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                                                                                                                            Yatapoxvirus
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                                                                                                                                                                                                                                                                                                                                                                                                                     Nakamura Y
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Brassicales;
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                                                                                                                  NCBI_TaxID=38804;
                                                                                                                                                                                                                                                                                                                                                                                                         Structural
                                                                                                                                                                                                                                                                   265
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itted (JAN-1999)
                                                                                                                                                                                                                                                                    GSLVTLSCVREEKL
                                                                                                                                                                                                                                                                                     GNLVTLSC--ETKL
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                                                                                                                                                                                                                                                                                                                                                         es. 7:131-135(2000).
AB022215; BAB01780.1; -
NCE 428 AA; 50007 MW;
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Similarity - 71.4%;
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                                                                                                                                                                                                                                                                                      12
           62.3%;
54.5%;
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                                                                             Yaba virus DNA.";
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Last annotation update)
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           Score 38;
Pred. No.
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Pred.
                                                EF74954D554EEBEF CRC64;
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4 bp cover
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                                                                                                                                        Poxviridae;
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dicots; Rosidae; eurosids II;
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                    Length
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Similarity 6; Conserv

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Q78242;
Q1-NOV-1996
Q1-NOV-1996
Q1-JUN-2000
                                                                                                                                                                                      Q9V539
Q9V539;
                                                                                                                                                                                                                                                                                                                                              PRINTS; PROUVER 192 AA; 22543 MW;
MEDLINE=20196006; PubMed=10731132; Adams M.D., Celniker S.E., Holt R.A., Amanatides P.G., Scherer S.E., Li P.W
                                                                               Drosophila melanogaster (Fruit fly).
Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Inse
Pterygota; Neoptera; Endopterygota; Diptera; Brachycera;
Ephydroidea; Drosophilidae; Drosophila.
                                                                                                                                         01-MAY-2000 (TremBLrèl.
01-MAY-2000 (TremBLrel.
01-OCT-2000 (TremBLrel.
BCDNA:GH08789 PROTEIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human immunodeficiency virus type 1.
Viruses; Retroid viruses; Retroviridae; Lentivirus.
NCBI_TaxID=11676;
                                     STRAIN-BERKELEY
                                              SEQUENCE FROM N.A.
                                                                                                                                BCDNA:GH08789
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      clones from
                                                                      NCBI_TaxID=7227;
                                                                                                                                                                                                                                                                                                                                                                                               INTERPRO; IPR000475; PFAM; PF00559; Vif; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                           Borsetti A., Saggio I.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Federico M., Titti F., Butto S., Orecchia A., Car
Macchi B., Mangiano N., Verani P., Rossi G.,
"Biologic and molecular characterization of produ
clones from HUT-78 cells infected with a patient
AIDS Res. Hum. Retroviruses 5:385-396(1989).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Carlini F., Federic M., Verani P., Rossi G.B.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                               1 GNLVTLSCE 9
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                                                                                                                                                                                                                                                                                                      Similarity 7; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                               Federico M., Butto S., Orecc
A., Saggio I., Verani P., Ros
mmunopharmacol. 3:17-23(1990)
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                                                                                                                                                                                                  PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                62.3%;
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                                                                                                                                                                                                 PRT;
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                                                                                                      Tracheata; Hexapoda; Insecta;
                                                                                                                                                                                                                                                                                                                                                                                                                                         Orecchia A., ., Rossi G.;
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 P.W.,
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 Evans C.A., Goca
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                                                                                                                                                                                                                                                                                                                           Length 192;
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 Gocayne J.D., A., Galle R.F.
                                                                                                                                                                                                                                                                                                      Indels
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isolate.";
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Q9Y3Y8:
01-NOV-1999 (TrEMBLrel. 12, Created)
01-NOV-1999 (TrEMBLrel. 12, Last sequence up
01-OCT-2000 (TrEMBLrel. 15, Last annotation
HYPOTHETICAL 63.9 KDA PROTEIN (FRAGMENT).
DKFZP56411922.
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Hostin D., Houston K.A., Howland T.J., Wei M.-H., Ibegwam C.,
Jalali M., Kalush F., Karpen G.H., Ke Z., Kennison J.A., Ketchum K.
Kimmel B.E., Ködira C.D., Kraft C., Kravitz S., Kulp D., Lai Z.,
Lasko P., Lei Y., Levitsky A.A., Li J., Li Z., Liang Y., Lin X.,
Liu X., Mattei B., McIntosh T.C., McLeod M.P., McPherson D.,
Merkulov G., Milshina N.V., Mobarry C., Morris J., Moshrefi A.,
Mount S.M., Moy M., Murphy B., Murphy L., Muzpy D.M., Nelson D.L.,
Nelson D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M.,
Palazzolo M., Pittman G.S., Pan S., Pollard J., Puri V., Reese M.G
Reinert K., Remington K., Saunders R.D.C., Scheeler S., Shen H.,
Chica B. C. (Schoel T. Stancer)
            Wambutt R., Heubner D., Mewes H.W., Gassenhuber J., Wiemann Submitted (MAY-1999) to the EMBL/GenBank/DDBJ databases. EMBL, AL049946; CAB43220.1; - HSSP; P56276; 1TLK
                                                                                                                                                         Eukaryota; Metazoa; Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Zheng X.H., Zhong F.N., Zhong W., Zhou X., Zhu S., Zhu Gibbs R.A., Myers E.W., Rubin G.M., Venter J.C.; "The genome sequence of Drosophila melanogaster.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Shue B.C., Siden-Kiamos I., Simpson M., Skupski M.P., Smith T., Spier E., Spradling A.C., Stapleton M., Strong R., Sun E., Svirskas R., Tector C., Turner R., Venter E., Wang A.H., Wang X., Wang Z.-Y., Wassarman D.A., Weinstock G.M., Weissenbach J., Williams S.M., Woodage T., Worley K.C., Wu D., Yang S., Yao Q.A., Ye J., Yeh R.-F., Zaveri J.S., Zhan M., Zhang G., Zhao Q., Zheng
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Beeson K.Y.,
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PFAM; PF02141;
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Sutton G.G.,
                                                                                   rissue=brain;
                                                                                                     SEQUENCE FROM N.A.
                                                                                                                                          NCBI_TaxID=9606;
                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Science 287:2185-2195(2000).
EMBL; AE003834; AAF58981.1;
FLYBASE; FB90027561; BCDNA:GH08789
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K.H., Doyle C., Baxter E.G., Helt G., Nelson C.
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letazoa; Chordata;
theria; Primates;
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Evangelista C.C., Ferraz C.
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50.0%;
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Pred. No.
                                                                                                                                                         Craniata; Vertebrata; Catarrhini; Hominidae;
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15;
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K., Zhu S., Zhu X.,
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Baldwin D.,
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INTERPRO; IPR003006; -

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584 AA; 63918 MW;

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078109;
01-NOV-1998 (TrEMBLrel. 08, C
01-NOV-1998 (TrEMBLrel. 08, L
01-JUN-2000 (TrEMBLrel. 14, L
Q9TPP1;
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01-MAY-2000 (
01-MAY-2000 (
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                                                                                                                                                                                                                                 Submitted (OCT-1998) to the EMBL; AF042159; AAC34367.1; EMBL; AF100710; AAF06839.1;
                                                                                                                                                                                                                                                                             Otten L.A., Steimle V., Bontron S., "Equivalent expression of the Major II genes through the various forms of the Major II.
                                                                                                                                                                                                                                                                      CIITA.
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                                               Q9TPP1
                                                                                                                                                                                                           SEQUENCE
                                                                                                                                                                                                                                                                                                                                        Submitted (JAN-1998) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                   STRAIN-BALB/C;
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"Expression of MHC class II molecules in different cellular functional compartments is controlled by differential usage multiple promoters of the transactivator CIITA.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRAIN-BALB/C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NCBI_TaxID=10090;
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117 GDVVTVACEAK 127
                                                                                                       798 GNLVGLSCVT
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Pred. No.
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r Histocompatibility Complex
of the mouse transactivator
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01-OCT-2000
01-OCT-2000
2361
                                                                                                                                                                                                         Crowl R.M., Luk D.;

'Identification of the gene encoding Adlican, a novel prexpressed in human arthritic tissues.";

Submitted (MAR-2000) to the EMBL/GenBank/DDBJ databases.

EMBL; AF245505; AAF86402.1;

EMBL; AF245505 AAF86402.1;

SEQUENCE 2828 AA; 312292 MW; A18377D8554F1FE1 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE-97327562; PubMed-9184229; Muhlethaler-Mottet A., Otten L.A., Steimle V., Mach B.; Muhlethaler mottet a., Otten L.A., Steimle V., Mach B.; Expression of MHC class II molecules in different cellular functional compartments is controlled by differential usage multiple promoters of the transactivator CITTA."; EMBO J. 16:2851-2860(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
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                                               1 GNLVTLSCETK
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  GDVVTVACEAK
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Last sequence update)
Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0,;
                                                                                                                        Score 38;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Craniata; Vertebrata; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Craniata; Yertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        F3FF05DEBFB0CE71 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    38;
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40;
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97;
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                                                                                                                                                                                                                                                                                                                         a novel protein
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                                                                                                                                                  Length 2828;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo
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                                                                                                Gaps
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Result
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Database
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OM protein -
                                                                                                                                                                                                                                                                                                  Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                   Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  seq
     55555555555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein search, using sw
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  length: 0
length: 20
                                                                                                                                                                                                                 Query
Match
                                                       1000
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Gapop 10.0 , Gapext 0.5
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61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          June
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 LLQRPGLQLYFS 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         390729 seqs, 57163235 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Geneseq_0401:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /SIDS6/gcgdata/geneseq/geneseqp/AA1984.DAT:/
/SIDS6/gcgdata/geneseq/geneseqp/AA1985.DAT:/
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                'SIDS6/gcgdata/geneseq/geneseqp/AA1986.DAT:
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 W80448
W97833
W97834
                                                                                       Y96183
R20811
R22550
                                                                                                                                         W60555
R22549
R91439
                                                       R91438
W00859
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /geneseq/geneseqp/AA1988.DAT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /geneseq,
                                                                                                                                                                                                                                                                     SUMMARIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /geneseqp/AA1989.DAT:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          390729
                                                                                                                                                                                                                 Description
                                                     Human FCRI (CDNA C
                  Human Fc receptor
                                                                                        Human macrophage-s
                                                                                                       Human macrophage-s
                                                                                                                                                                              Oligopeptide from
                                                                                                                                            Human FCRI (cDNA c
                                                                                                                                                               Human
Fc receptor
                                                                                                                                                         macrophage-s
```

Zea mays protein i	G12546	21	124	55.7	34	
an secreted pr	176	21	88	55.7	34	
	W02258	17	1411	57.4	35	
Protein containing	Y90279	21	9	57.4	. 35	
Arabidopsis thalia	G14485	21	365	57.4	35	
Arabidopsis thalia	G21246	21	9	57.4	35	
Arabidopsis thalia	G14486	21	9	57.4	35	
thal	G21247	21	249	57.4	35	
Arabidopsis thalia	G14487	21	249	57.4	<u>3</u> 5	
thal	G35900	21	201	57.4	35	
hal	G21248	21	201	57.4	35	
thal	G10127	21	201	57.4	3 5	
prot	Y90275	21	170	57.4	35	
· Arabidopsis thalia	G35901	21	129	57.4	35	
S.	G10128	21	129	57.4	35	
rain 17	W72975	20	838	59.0	36.	
ent of hum	Y41438	20	·319	59.0	36	
ൌ	Y73820	20	127	. 59.0	36	
Dengue virus type	W68417	19	401	60.7	37	
polyprotein.	R35081	14	3080	٠	38	
an smoothened	Y43263	20	803	٠	38	
3	B28645	22	793	•	38	
acid sequen	W81065	20	793	62.3	38 88	
smoothened	Y43262	20	793	•	38	
s norvegicu	W37911	19	793		38	
Smoot	B28646	22	, 787	٠	38	
o acid sequen	W81064	20	787	٠	38	
sapi	W37912	19	. 787		38	
FC r	W86195	20	377	86.9	Ś	
sfc-gammar	Y33183	20	261		55	
ma	R12428	12	410	91.8	56	
cancer ass	B43683	21	399	100.0	61	
high affi	Ý96226	21	374	100.0	61	
Human macrophage-s	Y96134	21	374	100.0	61	

ALIGNMENTS

PXX W60555 XX So 08-OCT-1996; 07-OCT-1997; 16-APR-1998. WO9815833-A1 Synthetic. cell transformation; mutation; anti; oncogene. Extracellular domain; CD64; identification; antibody; immunohistochemical; immunofluorescent analysis; detection; Oligopeptide from extracellular domain of CD64 18-AUG-1998 W60555 W60555 standard; peptide; 12 (UYUT-) RIJKSUNIV UTRECHT. (first entry) 96EP-0202791. 97WO-NL00557. B

Identifying peptide(s) binding specifically to protein target - by expressing on phage surface and testing for binding to immobilised oligo:peptide derived from the target, useful for, e.g. identifying specific antibodies

WPI; 1998-240964/21

De Kruif CA, Logtenberg T;

Example 1;

Page

29;

40pp; English

W60537-61

```
RESULT
R22549
                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 100.
Matches 12; Conservative
                            N-PSDB;
                                    WPI; 1992-056864/07
                                                     Seed B,
          New CD53 cell surface antigen
                                                                      (GEHO-)
                                                                                       13-JUL-1990;
                                                                                                                         23-JAN-1992
                                                                                                                                          WO9201049-A
                                                                                                                                                          Homo sapiens
                                                                                                                                                                         Rapid immunoselection cloning technique; cell surface antigen; immunodiagnosis; high affinity receptor.
                                                                                                                                                                                                  Human macrophage-specific FcRI receptor encoded by clone p98.
                                                                                                                                                                                                                     21-MAY-1992 (first entry)
                                                                                                                                                                                                                                         R22549;
                                                                                                                                                                                                                                                        R22549 standard; Protein;
                                                                                                       15-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                         .
2
                                                                                                                                                                                                                                                                                                  1 LLORPGLOLYFS 12
                                                                     GEN HOSPITAL
                                                    Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                               12 AA;
                                                                                      90US-0553759
                                                                                                       90WO-US04986
                                                     Amiot M;
                                                                                                                                                                                                                                                                                                                                           100.0%;
                                                                                                                                                                                                                                                        344 AA.
                                                                                                                                                                                                                                                                                                                                    0
of haematopoietic
                                                                                                                                                                                                                                                                                                                                           Score 61;
Pred. No.
                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                           9.5e-05;
                                                                                                                                                                                                                                                                                                                                                     BG
                                                                                                                                                                                                                                                                                                                                                     19;
                                                                                                                                                                                                                                                                                                                                   0;
it - for
neoplasms,
                                                                                                                                                                                                                                                                                                                                                  Length 12;
                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                   Gaps
```

This amino acid sequence was predicted from the cDNA sequence of cDNA clone p98/X2. It differs from the sequence predicted from clone p985 (see Q211/8) at position 25; p135 encodes a Ser residue while clone p98/X2 predicts a Thr residue. At position 58, p90

Ser residue 58, p90

of the p98 product

is truncated

compared

e; the C-terminal sequence with those of the p135

cDNA clone, p98 (T14718), obtd. from a cDNA library using an immunoselection cloning method. FcRI is a high affinity receptor for the FC portion of IgG, normally located on cell surfaces of macrophages. 2 Other cDNA clones, p135 (T14717) and p90 (T14719), coded for a variants of the FcRI sequence; the C-terminal sequence

The amino acid

sequence (R91439) of human

FcRI was detd.

Example 10;

Page 94a;

160pp;

English.

immuno-therapy

and diagnosis

```
RESULT
R91439
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  뫄
                                                                                                                                                                                                                                                                                 XX
AX
XX
Qd
                                                                                                                                                                                                                                                                                                                                                             FFF SXXXXXXX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
Matches 12; Conserv
                                                                                                                                                                                                                         01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
18-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (see Q21180) predicts Val and p98/X2 predicts Leu. Sequences predicted from all 3 clones show the typical features of a type I integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane-spanning domain, and a short, highly charged cytoplasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and six Cys residues distributed among three C2 set Ig-related domains. A fusion protein of FCRI and a receptor ligand will be helpful to increase the potency of antibodies in therapy.
                                                                                     Example 10; Column 55-56; 79pp; English.
                                                                                                                      Cloning of cDNA encoding cell surface antigen - useful for isolation
                                                                                                                                                                              Aruffo A,
                                                                                                                                                                                                                                                                                             25-FEB-1988;
                                                                                                                                                                                                                                                                                                                  09-APR-1996
                                                                                                                                                                                                                                                                                                                                        US5506126-A
                                                                                                                                                                                                                                                                                                                                                                                   Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                    Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human FCRI (cDNA clone p98 product).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      R91439;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           R91439 standard;
                                                                                                                                                                                                     (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                        therapy; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                diagnostic and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 LLQRPGLQLYFS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 w
                                                                                                                                               1996-200279/20
DB; T14718.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    surface
                                                                                                                                                                                                                                                                                                                                                                                                                   sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   344 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; ilarity 100.0%; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                               92US-0983647.
88US-0160416.
89US-0379076.
                                                                                                                                                                                                                          90US-0553759
93US-0139273
                                                                                                                                                                                                                                                                                           88US-0160416
                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers 58
                                                                                                                                                                                                                                                                                                                                                                        /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12
                                                                                                              therapeutic
                                                                                                                                                                                                                                                                                                                                                                                                                                        vector;
                                                                                                                                                                                                                                                                                                                                                                                                                                       cloning; immunoselection;
ector; FcRI; Fc receptor.
                                                                                                                                                                                                                                                                                                                                                           "amino acid 58 is Leu
translated product"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           344 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 61; DB 13;
Pred. No. 0.0027;
                                                                                                             proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                       in
                                                                                                                                                                                                                                                                                                                                                                       clone
                                                                                                                                                                                                                                                                                                                                                                                                                                                immunotherapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 344;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                     p135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
```

0;

epitopes on different molecules. The genes/oligonucleotides that encode selected peptides can be isolated and used for recombinant production of antibodies. Selected antibodies are used, e.g. for immunohistochemical or immunofluorescent analysis, and also to detect cell transformation caused

0;

by mutation

in (anti)oncogenes.

surface of a replicable display package, synthesising oligopeptides derived from the target protein on a solid phase, and testing for binding between the peptide and oligopeptides. The method is used to screen large peptide libraries, especially to detect antibodies, or their fragments, that bind to cell markers or that can differentiate between different forms of the same protein, including bispecific antibodies that bind to two non-overlapping epitopes on the same monomeric antigen or two

Synthetic oligopeptides W60537-61 are derived from the extracellular domain of CD64. They were synthesised on 25 polyethylene rods as adjacent, non-overlapping sets. Peptide specific phages were isolated by affinity binding to the CD64-covered rods. The specification describes a method for the identification of a peptide able to bind specifically to a target protein. The method comprises displaying the peptide on the curfect of a requirement.

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RESULT
Y96183
ID Y9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SSSSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 and p90 products (see also R91438 and W00859). A fusion protein of FCRI and a receptor ligand will be helpful in increasing the potency of antibodies in therapy.
                                                                                                                                                                                                                                                                                         /note=
Misc-difference 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 panning; immunodiagnosis; diagnosis; immunotherapy; gene therapy; immune disorder; infection; asthma; immune-complex disease; amyloidosis; multiple sclerosis; parasitic disease; Fc receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
Misc-difference
                     Misc-difference
                                          Misc-difference 139
                                                           /note=
Misc-difference 136
                                                                                          Misc-difference
                                                                                                    /note= "encoded
Misc-difference 129
                                                                                                                                       /note=
Misc-difference 126
                                                                                                                                                                                     Misc-difference
                                                                                                                                                                                                          Misc-difference
                                                                                                                                                                                                                            /note=
Misc-difference 116
                                                                                                                                                                                                                                                       Misc-difference
                                                                                                                                                                                                                                                                                                                         Misc-difference
                                                                                                                                                                                                                                                                                                                                                 Misc-difference
                                                                                                                                                                                                                                                                                                                                                                        Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                               Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                     Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human macrophage-specific FcRI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-DEC-2000 (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Y96183;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Y96183 standard; Protein; 344 AA
                                                                                                                                                               Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Macrophage; FcRI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          217
                                                                                                                                                                                                                                                                                                                                                                                                                                            Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                y Match 100.0%;
Local Similarity 100.0%;
nes 12; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 LLQRPGLQLYFS 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       llqrpglqlyfs 228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              344 AA;
                                                                                                                                                                                                                                                   /note=
                                                                                                                                                                                                                                                                                                                                                                                                                /note=
. 38
                                                                                          /note=
134
                                                                                                                                                              /note-
123
                                                                                                                                                                                     /note=
122
                                                                                                                                                                                                          /note=
117
                                                                                                                                                                                                                                                                                                                                                 /note-
55
                     /note=
140
                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                        /note=
                                                                                                                                                                                                                                                                                                                                     'note=
                                                                                                                                                                                                                                                                                                                                                                                  'note=
                                                                                                                                                                                                                                                                                                                                                                                                         'note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cell surface antigen;
                                                                                                                                                   "encoded
                                                                                                                                                                                                                                                                                                                                   "encoded by CAC"
                                                                                                                                                                                                                      "encoded
                                                                                                                                                                                                                                                                                      "encoded by CCC"
                                                                                                                                                                                                                                                                                                                                                                                  "encoded by CGG"
                                                                                                                                                                                                                                                                                                                                                                                                                               "encoded by GCG"
         "encoded by AAT"
                                "encoded
                                                       "encoded
                                                                                                                                                                        "encoded
                                                                                                                                                                                               "encoded
                                                                                                                                                                                                                                            "encoded
                                                                                                                                                                                                                                                                  "encoded by CAG"
                                                                                                                                                                                                                                                                                                             "encoded by
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                                                                                                                                                                                                                                                                                                                                                                                                        "encoded by ACC"
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Pred. No. 0.0027;
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                                                                                                                                                                                                                    by GAA"
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                                                       þу
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                                                                                                                                                                       by TTC"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                 CCA"
                                                       GTT"
                                                                                                                            cce"
                                                                                                                                                                                                                                            CAA"
                                                                                                                                                                                                                                                                                                                                                           ACC "
                                                                              AAT"
                                                                                                                                                   ACC"
                                                                                                                                                                                               GCA"
                                                                                                                                                                                                                                                                                                                TCC"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     immunoselection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 344;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
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1 LLQRPGLQLYFS 12

Query Match Best Local S Matches 12

Local Similarity

100.0%;

Score 61; Pred. No.

0.0027

DB 21; 0

Length 344; Indels

0

Gaps

0;

0;

Mismatches

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13-JUL-1989;
23-MAR-1990;
13-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note-
Misc-difference 332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Misc-difference
                                                                                                                                                          immunotherapeutic applications, including the diagnosis and treatment of immune-mediated infections, diseases, and disorders in animals, including humans. These disorders include asthma,
                                                                                                                                                                                                                                                                                                                        The present sequence is that of a human macrophage specific FCRI, as deduced from cDNA clone pl35 (see A50631), which was isolated from a cDNA library expressed in COS cells using a novel method of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Stamenkovic I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Misc-difference
Sequence
                                                      protein of
                                                                       surfaces other than macrophages,
                                                                                                                            sclerosis.
                                                                                                                                                                                                                   CSA nucleic acids isolated by the method of the invention, and proteins they encode, are useful for immunodiagnostic and
                                                                                                                                                                                                                                                                         physica.
                                                                                                                                                                                                                                                                                          based upon transient expression of a CSA in eukaryotic cells and
                                                                                                                                                                                                                                                                                                           the invention designed to isolate CSA nucleic acids.
                                                                                                                                                                                                                                                                                                                                                                                                  Example 10; Column 53-55; 75pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Isolated nucleic acid molecule encoding the CD19 cell surface antigen, useful for immunodiagnosis and immunotherapy of immune-mediated infections or disorders, e.g. asthma, immune-complex disease, parasiti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-DEC-1992;
25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US6111093-A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Misc-difference
                                                                                        ability to interfere with such bonding,
                                                                                                                          immune-complex disease, amyloidosis, parasitic diseases or multiple sclerosis. FCRI is a high affinity receptor for the Fc portion of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                      panning on) an antibody-coated substrate such as a culture
                                                                                                         rosis. FCRI is a high affinity receptor normally located on the cell surfaces
                                                                                                                                                                                                                                                                       selection of cells expressing the antigen by adhesion
   344
                                                      FcRI and
 ĀĀ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       89US-0379076.
90US-0498809.
90US-0553759.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Seed B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                92US-0983647
88US-0160416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    98US-0181612
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333
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306
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305
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220
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216
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note=
                                                      a receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "encoded by GTG"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "encoded
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "encoded by
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "encoded by CAA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "encoded by
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                                                      ligand will
                                                                                                                                                                                                                                                                                                                                                                                                  English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GGC"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAC"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAT"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CGT"
                                                    ces of macrophages. The or to cause it to occur on ful in therapy. A fusion ill be helpful to increase
                                                                                                                                                                                                                                                                                                           The method
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217 llgrpglqlyfs 228

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                                                                                                                               RESULT
                                                                                                                                                                                                                                                                                                                                                                                    Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 12; Conservative C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This amino acid sequence was predicted from the cDNA sequence of cDNA clone p135. It differs from the sequence predicted from clones p98/X2 and p90 (see Q21179 and Q21180, respectively) at position 25; p135 encodes a Ser residue and the other two clones predict a Thr residue. At position 58, p135 predicts Leu and p90 predicts Val. Sequences predicted from all 3 clones show the typical features of a type I integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane-spanning domain, and a short, highly charged membrane-spanning domain, and a short, highly charged
                                                                                                                                                                                                                                                                                                                                                                                                                   cytoplasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and six Cys residues distributed among three C2 set Ig-related domains. A fusion protein of FCRI and a receptor ligand will be helpful to increase the potency of antibodies in therapy.
                                           R22550;
                                                                                 R22550 standard;
21-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                   1 LLQRPGLQLYFS 12
||||||||||
| 217 llqrpglqlyfs 228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 10; Page 94a; 160pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New CD53 cell surface antigen and DNA encoding it for immuno-therapy and diagnosis of haematopoietic neoplasms, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Seed B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23-JAN-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      W09201049-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human macrophage-specific FcRI receptor encoded by clone pl35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21-MAY-1992 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R20811;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R20811 standard; Protein; 374 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (GEHO-) GEN HOSPITAL CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rapid immunoselection cloning technique; cell surface antigen; immunodiagnosis; high affinity receptor.
                                                                                                                            6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                374 AA
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           90US-0553759
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    90WO-US04986
                                                                                   Protein; 374
                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                         Score 61; DB 13;
Pred. No. 0.003;
); Mismatches 0;
                                                                                   A.
                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                Length 374;
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Query Match
Best Local Similarity
Thes 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         domains. A helpful to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            and clone p90 predicts a Thr residue. At position 58, p90 predicts val while clones p135 and p98/XZ (see Q21179) both predict Leu. Sequences predicted from all 3 clones show the typical features of a type I integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane spanning domain, and a short, highly charged cytoplasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and six Cys residues distributed among three C2 set Ig-relate domains. A fusion protein of FCRI and a receptor ligand will be beliefed.
                                              Key
Misc-difference
                                                                                                                                                                  Cell surface antigen; cloning; immunoselection; therapy; diagnosis; vector; FcRI; Fc receptor.
                                                                                                                                                                                                                                  Human FcRI (cDNA clone pl35 product).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This amino acid sequence was predicted from the cDNA sequence of cDNA clone p90. It differs from the sequence predicted from clone p135 (see Q21178) at position 25; p135 encodes a Ser residue
                                                                                                                      Homo
                                                                                                                                                                                                                                                                                      30-OCT-1996
                                                                                                                                                                                                                                                                                                                                     R91438;
                                                                                                                                                                                                                                                                                                                                                                                 R91438 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 10; Page 94a; 160pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New CD53 cell surface antigen and DNA encoding it - for immuno-therapy and diagnosis of haematopoietic neoplasms, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1992-056864/07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Seed B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         W09201049-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rapid immunoselection cloning technique; cell surface antigen; immunodiagnosis; high affinity receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          23-JAN-1992
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (GEHO-) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             217 llgrpglglyfs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 LLQRPGLQLYFS
                                                                                                                   sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           fusion protein of FCRI and a receptor increase the potency of antibodies in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%;
Conservative (
                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             90US-0553759
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                                            Location/Qualifiers 25
/note= "amino acid 25 is Thr in clone p90 and p98
    translated products"
                                                                                                                                                                                                                                                                                                                                                                               Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     228
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                                                                                                                                                                                                                                                                                                                                                                                 374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 61;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 13; Length 374; 0.003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0
                                                                                                                                                                                      immunotherapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                glycosylation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
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Misc-difference 58

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Query Match
Best Local :
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01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The amino acid sequence (R91438) of human FCRI was detd. from a CDNA clone, pl35 (T14717), obtd. from a cDNA library using an immunoselection cloning method. FCRI is a high affinity receptor
                                                                                                                                                                                                         therapy; diagnosis;
                                                                                                                                                                                                                       Cell surface antigen; cloning; immunoselection;
                                                                                                                                                                                                                                                                                                                               W00859;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cloning of cDNA encoding cell surface antigen - useful of diagnostic and therapeutic proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-DEC-1992;
25-FEB-1988;
                                                                                                    09-APR-1996
                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                               30-OCT-1996
                                                                                                                                                                                                                                                                                                                                                               W00859 standard; Protein; 374 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           for the Fc portion of IgG, normally located on cell surfaces of macrophages. Another isolated cDNA clone, p90 (T14719), coded for a variant (WO0859) of the FcRI sequence, and a third clone, p98 (T14718), coded for an FcRI (R91439) having a different C-terminal sequence. A fusion protein of FcRI and a receptor ligand will
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1996-200279/20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     09-APR-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US5506126-A
                                                                    25-FEB-1988
                                                                                                                                      US5506126-A
                                                                                                                                                                                                                                                         Human FcRI (cDNA clone p90 product).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 10; Column 55-56; 79pp; English:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18-OCT-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-FEB-1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                       217
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13-JUL-1989;
13-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity 100 hes 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 LLQRPGLQLYFS 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    llqrpglqlyfs 228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Seed B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             in increasing the potency of antibodies in therapy.
                                                                                                                                                                                                                                                                                           (first entry)
92US-0983647.
88US-0160416.
89US-0379076.
                                                                    88US-0160416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               93US-0139273
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900S-0553759
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               92US-0983647.
88US-0160416.
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                                                                                                                                                                                                          vector; FcRI; Fc receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "amino acid 58 is Val in p90 clone
translated product"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 61; DB 17
Pred. No. 0.003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                         immunotherapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 374;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
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33
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W80448
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Matches 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match .
Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        for the FC portion of IgG, normally located on cell surfaces of macrophages. Other CDNA clones (see also T14717-18) coded for variants (R91438-39) of the FCRI sequence. A fusion protein of FCRI and a receptor ligand will be helpful in increasing the potency of antibodies in therapy.
                      Misc-difference
                                            Misc-difference
                                                                   Misc-difference
                                                                                          Misc-difference
                                                                                                                                         Misc-difference
                                                                                                                                                               Misc-difference
                                                                                                                                                                                      Misc-difference
                                                                                                                                                                                                                                                                                              cloning
                                                                                                                                                                                                                                                                                                                                 Human Fc
                                                                                                                                                                                                                                                                                                                                                        07-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                    W80448 standard; Protein; 374 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The amino acid sequence (W00859) of human FCRI was detd. cDNA clone, p90 (T14719), obtd. from a cDNA library using immunoselection cloning method. FCRI is a high affinity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cloning of cDNA encoding cell surface antigen - useful for isolation of diagnostic and therapeutic proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           N-PSDB; T14719
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1996-200279/20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13-JUL-1990;
18-OCT-1993;
                                                                                                                                                                                                            Misc-difference
                                                                                                                                                                                                                                     Misc-difference
                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                        Fc receptor I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (GEHO ) GEN
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                                                                                                                                                                                                                                                                                                                              receptor I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Column
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HOSPITAL CORP
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                                                                                                                /note=
. 77
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93US-0139273.
/note=
159
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60
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                       141
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23
                                             103
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                                                                                                                                                                                                                                            Location/Qualifiers
                                                                              /note=
                                  'note=
                                                         note=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   55-56; 79pp; English.
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                                                                                                                                                                                                                                                                                                        cell surface antigen; lymphocyte; human;
                                                                                                   "encoded by AAT"
                                                                                                                                                 "encoded by
           "encoded by
                                                                                                                             "encoded
                                  "encoded
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                                                                              "encoded
                                                                                                                                                                         "encoded by GAC"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 61;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   from a cDNA library using an FCRI is a high affinity receptor
           GGC
                                                                              TCC.
                                  222
                                                         CAA
                                                                                                                             CCC"
                                                                                                                                                    CTG"
                                                                                                                                                                                                  CTC"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 374;
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W97833
                                                      RESULT 10
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Best Local
                                                                                                                                                 Matches
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13-JUL-1989;
13-JUL-1990;
                                                                                                                                                                                                                                                                 produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V63444) which allow the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and immunotherapeutic applications, including the diagnosis and
                                                                                                                                                                                                                                                                                                                                               expression libraries is based on transient expression of an antigen in eukaryotic cells and selection of cells expressing the antigen by adhesion to an antibody-coated substrate. The method is useful for the isolation and cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V63442-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated
  W97833;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New cloning vector and polylinker - based on efficient cloning and expression of mammalian human lymphocyte antigenic sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Misc-difference
                           W97833 standard;
                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                       genes can be expressed in a prokaryotic or eukaryotic host cells to produce the encoded protein. The invention also provides high
                                                                                                                                                                                                                                                                                                                                                                                                                                             from a single patient undergoing extracorporeal interlinduction therapy. Clones p90 (see X07372) and p98 (swere also obtained. A novel method for cloning cDNAs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This is the amino acid sequence of human Fc receptor I (FCRI), deduced from cDNA clone pl35 (see V63456) isolated using a rapi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      N-PSDB;
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                                                                                                                                                                                                                                                       immunotherapeutic applications, including the diag treatment of immune-mediated infections, diseases,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 10;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-MAY-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  immunoselection cloning method from a cDNA library expressed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (GEHO)
                                                                                             217
                                                                                                                                                            Local
                                                                                          llqrpglqlyfs
                                                                                                                      LLQRPGLQLYFS 12
                                                                                                                                                 1 Similarity
12; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GEN HOSPITAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The cDNA library was constructed from polyA RNA of cells
                                                                                                                                                                                                                                          including
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Column
                                                                                                                                                100.0%;
llarity 100.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              92US-0983647.
88US-0160416.
89US-0379076.
90US-0553759.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        256
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176
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171
                          Protein;
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                                                                                             228
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                                                                                                                                                                                                                                            humans.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "encoded
                             374 AA
                                                                                                                                                 0;
                                                                                                                                                            Score 61; DB 19;
Pred. No. 0.003;
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                                                                                                                                                 Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          existing sequences n cDNA(s), especiall
                                                                                                                                                                                                                                                                                                                                                                                                                                                          p98 (see X07373)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         interleukin-
                                                                                                                                                                       Length 374;
                                                                                                                                                 Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                              from mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sequences for especially
                                                                                                                                                                                                                                                        disorders
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                                                                                                                                                Gaps
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  FEFFF
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25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
21-MAY-1997;
induction therapy. were also obtained. expression libraries
                                                                                                                                              New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially human lymphocyte antigenic sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note=
Misc-difference 159
            from a single patient undergoing extracorporeal interleukin-2 induction therapy. Clones pl35 (see V63456) and p98 (see X07373) were also obtained. A novel method for cloning cDNAs from mammal
                                                     This is the amino acid sequence of human Fc receptor I (FCRI), as deduced from cDNA clone p90 (see X07372) isolated using a rapid immunoselection cloning method from a cDNA library expressed in COS cells. The cDNA library was constructed from polyA RNA of cells
                                                                                                                                                                                                                                          Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                       03-NOV-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human Fc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-JUN-1999
                                                                                                                      Example 10; Column 53-54; 75pp;
                                                                                                                                                                                                                                                                                                                                                                              21-MAY-1997;
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                                                                                                                                                                                                                                                                    (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                   1998-609251/51.
DB; x07372.
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                                                                                                                                                                                                                                            Seed
    libraries
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                                                                                                                                                                                                                                                                                                           88US-0160416.
89US-0379076.
90US-0553759.
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77
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23
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176
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141
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                                                                                                                                                                                                                                            В;
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171
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    is based on
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     genes can be expressed in a prokaryotic or eukaryotic host cells to produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V63444) which allow the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and immunotherapeutic applications, including the diagnosis and treatment of immunotherapeutic applications, including the diagnosis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       in eukaryotic cells and selection of cells expressing the antigen by adhesion to an antibody-coated substrate. The method is useful for the isolation and cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V63442-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated
                                                                                       /note=
Misc-difference 171
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US5830731-A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Fc receptor I; FcRI; cell surface antigen; lymphocyte; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human Fc receptor I
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217 llqrpglqlyfs 228
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ilarity 100.0%;
Conservative (
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159
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256
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PRPRXTX
                                                                        induction therapy. Clones plass (see V63456) and p90 (see X07372)

Covere also obtained. A novel method for cloning cDNAs from mammalian expression libraries is based on transient expression of an antigen of in eukaryotic cells and selection of cells expressing the antigen by adhesion to an antibody-coated substrate. The method is useful for the isolation and cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V6344-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated genes can be expressed in a prokaryotic or eukaryotic host cells to produce the encoded protein. The invention also provides high the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and immunotherapeutic applications, including the diagnostis and treating at immunocharapeutic applications, including the diagnostis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13-JUL-1989;
13-JUL-1990;
21-MAY-1997;
Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                     This is the amino acid sequence of human Fc receptor I (FCRI), as deduced from cDNA clone p98/X2 (see X07373) isolated using a rap immunoselection cloning method from a cDNA library expressed in cells. The cDNA library was constructed from polyA RNA of cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially human lymphocyte antigenic sequences
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25-FEB-1988;
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                                                               treatment of immune-mediated infections,
                                                                                                                                                                                                                                                                                                                                                                            from a single patient undergoing extracorporeal interleukin-2 induction therapy. Clones pl35 (see V63456) and p90 (see X07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 10; Column 53-54; 75pp; English.
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DB; x07373.
                                       including
  374 AA;
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90US-0553759.
97US-0861205.
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880S-0160416
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Query Match
Best Local Similarity
Matches 12; Conser 1 LLQRPGLQLYFS 12 Conservative 100.0%; 0, Score 61; DB 19 Pred. No. 0.003; Mismatches DB 19; Length 374; Indels 0, Gaps

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RESULT Y96134 19-DEC-2000 Y96134; Human macrophage-specific FcRI. Y96134 standard; Protein; 374 AA. (first entry)

XSXEX

amyloidosis; Macrophage; FcRI; cell surface antigen; human; immunoselection; immune panning; disorder; immunodiagnosis; diagnosis; infection; asthma; sis; immunotherapy; gene th
ma; immune-complex disease;
parasitic disease; Fc rece therapy;

2000-586382/55 DB; A50592.

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01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
23-MAR-1990;
13-JUL-1990;
Stamenkovic I,
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                (GEHO ) GEN HOSPITAL CORP
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Seed B;

WPI; 2000-367968/32 N-PSDB; A27466.

(PLAC) MAX PLANCK GES FOERDERUNG WISSENSCHAFTEN

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RESULT
Y96226
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Best Local
                                                                                                                                                                                 Human; high affinity Fc receptor; FcgammaRI; immunoglobulin; infection; immune response; CD64; monocyte; macrophage; neutrophil; eosinophil; HTV; 196; immunosuppressive; antirheumatic; cytostatic; antiinflammatory; AIDS; SLE; multiple myeloma; rheumatoid arthritis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ability to interfere with such bonding, or
surfaces other than macrophages, is useful
protein of FCRI and a receptor ligand will
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CSA nucleic acids isolated by the method of the invention, and the proteins they encode, are useful for immunodiagnostic and immunotherapeutic applications, including the diagnosis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present sequence is that of a human macrophage specific FCRI, as deduced from cDNA clone pl35 (see A50592), which was isolated from a cDNA library expressed in COS cells using a novel method of
03-DEC-1998;
                                 03-DEC-1998;
                                                                07-JUN-2000.
                                                                                                EP1006183-A1
                                                                                                                                Homo sapiens
                                                                                                                                                                   systemic lupus
                                                                                                                                                                                                                                                                   Human high affinity Fc receptor,
                                                                                                                                                                                                                                                                                                      11-SEP-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          immune-complex disease, amyloidosis, parasitic diseases or multiple sclerosis. FCRI is a high affinity receptor for the FC portion of IGG, normally located on the cell surfaces of macrophages. The ability to interfere with such bonding, or to cause it to occur on surfaces other than macrophages, is useful in therapy. A fusion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           treatment of immune-mediated infections, diseases, and diso
animals, including humans. These disorders include asthma,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   physical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the invention designed to isolate CSA nucleic acids.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           217
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              physical selection of cells expressing the antigen by adhesion panning on) an antibody-coated substrate such as a culture dis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 LLQRPGLQLYFS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             potencies of antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                        upon transient expression of a CSA in eukaryotic cells and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           l Similarity
12; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Column 53-55; 75pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                    (first entry)
98EP-0122969
                                 98EP-0122969
                                                                                                                                                                   erythematosus; tumour.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             in therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 61; DB 21
Pred. No. 0.003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 English.
                                                                                                                                                                                                                                                                   FcgammaRI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             diseases, and disorders in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           in therapy.
be helpful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 374;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              to increase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The method
                                                                                                                                                                                    arthritis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  parasitic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antigen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           may be modified to produce recombinant versions. The recombinant FC receptor consist only of the extracellular portion of the receptor and are not glycosylated; e. they do not have transmembrane domains or signal peptides. The recombinant proteins may be used in immunoassays to determine the immune status of patients with chronic diseases of the immune system, e.g. AIDS, systemic lupus erythematosus (SLE), multiple myeloma (MM), or rheumatoid arthritis. In addition, pharmaceutical compositions containing recombinant proteins may be used to treat or prevent autoimmune diseases, allergies or tumours, especially AIDS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     role in defending the body against infections. First, pathogens are opsonised by serum immunoglobulins. The resulting complex then binds cells expressing for receptors. FogammaRI molecules are expressed by monocytes and macrophages, but expression can also be induced on neutrophils and eosiophils. Upon Fc receptor activation, immune efformet trophils and eosiophils.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 allergic reaction; graft versus host disease; organ rejection; haemostatic; thrombolytic; cardiovascular disorder; infection:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence is the human high affinity Fc receptor, FcgammaRI FcgammaRI is also known as CD64. Fc receptors play an important
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antidiabetic; antiasthmatic; antirheumatic; antiarthritic; antiviral;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   B43683
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                B43683 standard; Protein; 399
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            neutrophils and eosinophils. Upon Fc receptor activation, immune effector pathways are activated, leading to immune response. The present sequence may be modified to account the control of the control 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 26-28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    glycosylation,
cancer -
WPI; 2000-587533/55
                                                                                                                                                                                                                                                                                                                                                                                                   08-MAR-2000; 2000WO-US05882
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200055350-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          dermatological; neuroprotective; thrombolytic; coagulant; nootropic;
vasotropic; antipsoriatic; antiangiogenic; gene therapy; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  antiinflammatory; antithyroid; antiallergic; antibacterial; cardiant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; cancer associated gene; cancer antigen; detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human cancer associated protein sequence SEQ ID NO:1128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 08-FEB-2001
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                                                                                                       Rosen CA,
                                                                                                                                                                                              (HUMA-) HUMAN GENOME
                                                                                                                                                                                                                                                                                                 12-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
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||||||||||||
||17 llqrpglqlyfs 228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            l Similarity
12; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 arthritis
                                                                                                       Ruben SM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                 99US-0124270.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%;
                                                                                                                                                                                                    SCI INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         proliferative; vulnerary;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   60pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ₹
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 61;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.003;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
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g immune disorders and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    immunomodulator;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cancer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
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WPI; 1991-164135/22

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Best Local Similarity
Thes 12; Conserv
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                                                                                                             PXPX
                                                                                                                                                                                                                                                                                                                      FΗ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             polypeptides, antibodies, agonists and antagonists from the present invention may be used to treat immune disorders by activating or inhibiting the proliferation, differentiation or mobilisation of immune cells, to treat disorders of haematopoietic cells, autoimmune disorders, allergic reactions, graft versus host disease and organ rejection, modulate haemostatic or thrombolytic activity, modulate inflammation, cancers, cardiovascular disorders, neurological disease and bacterial or viral infections. The peptides, nucleotides, antibodies, agonists and antagonists may be also be used in drug screens. C78449 to C78457 and B44240 represent sequences used in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     neuroprotective; cardiant; thrombolytic; coaqulant; nootropic; vasotropic; antipsoriatic and antiangiogenic. The polynucleotides and polypeptides can be used for preventing, treating or ameliorating med conditions and diagnosing pathological conditions. Polynucleotides,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C77607 to C78448 encode the human cancer associated proteins B43398 to B44239. The proteins can have activities based on and cells the genes are expressed in Example of activities
                                                                                                                                                                                                                                                                                                                                                                                            Hybrid Fc(gamma)RII/I receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                   14-AUG-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          R12428
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         R12428 standard; Protein; 410
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   antithyroid; antiallergic; antibacterial; antiviral; dermatological;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antiasthmatic; antirheumatic; antiarthritic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cytostatic;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel isolated nucleic acids comprising sequences encoding peptides useful for treating or diagnosing e.g.\ 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               N-PSDB; C77892
Hogarth PM,
                                                                        25-OCT-1989;
                                                                                                                                                  16-MAY-1991
                                                                                                                                                                                       W09106570-A
                                                                                                                                                                                                                                                               Region
                                                                                                                                                                                                                                                                                                                                                       Immunoglobulin-binding molecule; Fc(gamma)RI; Fc(gamma)RII
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                invention
                                   (UYME-) UNIV MELBOURNE
                                                                                                             25-OCT-1990;
                                                                                                                                                                                                                                                                                                  Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       242 llgrpglglyfs 253
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        proliferative; vulnerary; immunomodulator; antidiabetic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
Hulett MD,
                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                        89AU-0007045
                                                                                                               90WO-AU00513
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         encode the human cancer associated proteins given in . The proteins can have activities based on the tissues
                                                                                                                                                                                                                                                                 204..410
                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                              /label=
                                                                                                                                                                                                                                                                               /label= Fc(gamma)RII domains dl and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%;
                                                                                                                                                                                                                                            Fc(gamma)RI
 Ierino FL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 61;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    English
                                                                                                                                                                                                                                            D3,
 McKenzie IFC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; DB 21;
                                                                                                                                                                                                                                            transmembrane and
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DR N-PSDB; Q12079.

XX

New immunoglobulin binding hybrid Fc receptor molecules - used to PT control auto-immune diseases and allergic reactions and to regulate PT antibody prodn.

XX

Claim 10; fig 3B; 80pp; English.

XX

Claim 10; fig 3B; 80pp; English.

XX

Claim 10; fig 3B; 80pp; English.

XX

CC

This chimeric sequence is one example of a hybrid Fc receptor CC

CC functions of the constituent receptor types. It can be used for CC

CC requiating antibody production in vivo and in plasmaphoresis to CC

CC remove immune complexes or pathological antibodies. Rabbit IgG

CC bound to cells transfected with the chimeric Fc(gamma) receptor.

XX

XX

Sequence 410 AA;

Query Match 91:8%; Score 56; DB 12; Length 410;

Best Local Similarity 91:7%; Pred. No. 0.026;

Matches 11; Conservative 1; Mismatches 0; Indels 0; Gaps

Qy 1 LLQRPGLQLYFS 12

Db 232 llqrpglqlhfs 243

Search completed: June 4, 2001, 12:13:28

Search completed: June 4, 2001, 12:13:28
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Total number of hits satisfying chosen parameters:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score
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Copyright (c) 1993 - 2000 Compugen Ltd
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/cgn2_6/ptodata/2/iaa/PCTUS_COMB.pep:*
/cgn2_6/ptodata/2/iaa/backfiles1.pep:*
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PCT-US95-13703-21
US-09-293-395-2
US-09-293-395-5
US-09-293-395-13
US-09-293-395-13
US-09-053-197A-33
US-07-857-224B-48
US-07-857-224B-47
US-08-630-916A-46
                                                                                                                                                                                                                   US-08-284-393B-12
PCT-US95-08950-12
US-07-952-853-22
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US-08-840-316-3
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	Sequence 4, Appli	Sequence 5, Appli	Sequence 5, Appli	Sequence 9, Appli	Sequence 9, Appli	Sequence 2, Appli	Sequence 2, Appli	Sequence 9, Appli	Sequence 15, Appl	Sequence 15, Appl	Sequence 2, Appli	Sequence 19, Appl	Sequence 19, Appl	Sequence 7, Appli	Sequence 11, Appl	Sequence 8, Appli	Sequence 48, Appl	Sequence 4, Appli

ALIGNMENTS

US-08-332-562A-94

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TOPOLOGY:
US-08-332-562A-94
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                                                                                                                                                                                                                                                    PILING DATA:
CLASSIFICATION: 435
CLASSIFICATION DATA:
PRIOR APPLICATION NUMBER: US 07/
APPLICATION NUMBER: 1992
                                                                                                     TELEX: 904136
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Rel CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
                                                                                                                                    TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
TELEFAX: (202)672-5399
                                                                                                                                                                                         FILING DATE: 27-MAY-1992
ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 54270/119/GRHA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: HULETT, Mark D. TITLE OF INVENTION: FC RECEPTOR FOR IMMUNOGLOBULIN
                                                                                    SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
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                                                                  LENGTH:
                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/O
FILING DATE: 31-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
                                  STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE:
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20007-5109
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                                               : 91 amino acids
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                  ·linear
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                                 single
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                                                                                                       94:
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Best Local Similarity

91.7%;

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JS-08-953-823A-4
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US-08-720-484A-4
                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                     Sequence 4, Application US/08953823A Patent No. 6136958
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 4, Application US/08720484A Patent No. 5990281
                                                                                                                                                                      ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO:
                                                                                               APPLICANT: DeSauvage; Frederic
APPLICANT: Rosenthal, Arnon
APPLICANT: Stone, Donna
APPLICANT: Stone, Donna
TITLE OF INVENTION: Vertebrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 787 amino acids
                                                                 CORRESPONDENCE ADDRESS:
                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                      579 LLQNPGQELSFS 590
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb |
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: WinPatin (Gen
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: DeSauvage, Fi
APPLICANT: Rosenthal, Ai
APPLICANT: Stone, Donna
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                          Local Similarity
nes 8; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: FILING DATE: 30-Sel CLASSIFICATION: 53
                                 STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Syoboda, Craig G. REGISTRATION NUMBER: 39,044 REFERENCE/DOCKET NUMBER: P1
                                                   ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY:
                                                                                                                                                                                                                                                                                                                        1 LLQRPGLQLYFS 12
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South San Francisco
California
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                               E: Genentech, Inc.
1 DNA Way
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N: 530
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Pred. No. 41;
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1; Mismatches 0;
                                                                                                 Smoothened Proteins
                                                                                                                                                                                                                                                                                                                                                          Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                      Length 787;
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US-08-720-484A-2
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Best Local Similarity 66.7%;
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Patent No. 5990281
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 787 amino acids
TYPE: Amino Acid
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1489
TELEFAX: 650/952-9881
                                                                       ATTORNEY/AGENT INFORMATION:
NAME: Svoboda, Craig G.
REGISTRATION NUMBER: 39,044
                                                                                                                                                                                                                                                               COMPUTER READABLE FORM: mEDIUM TYPE: 3.5 inch, 1.44 mb floppy disk
                                                                                                                                                     SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION UNMER: US/08/720,484A
FILING DATE: 30-Sep-1996
                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: !
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION: TELEPHONE: 650/225-1489
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: WinPatin (GOURRENT APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/0:
FILING DATE: 30-Sep-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                           STREET: 1 DNA TW. CITY: South San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         579 LLQNPGQELSFS 590
                                                         REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                               COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                   CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC
OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: FILING DATE: 30-Sep CLASSIFICATION: 424
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 39, REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 LLQRPGLQLYFS 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /AGENT INC...
Syoboda, Craig G.
Syoboda, Craig G.
Syoboda, Craig G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Application US/08720484A
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Stone, Donna
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DeSauvage, Frederic
                                                                                                                                                                                                                                                                                                                                                                                                     Genentech,
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30-Sep-1997
N. /
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: 793 amino acids Amino Acid

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Query Match
Best Local Similarity
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                                                  US-08-216-260-4
                                                                RESULT
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Sequence 4, Application US/08216260 Patent No. 5837261 GENERAL INFORMATION:
                                                                                                                                                                      Query Match
Best Local Similarity Matches 8; Conserva
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                                                                                                                                                                                                                                                                                                                         NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                NAME: Svoboda, Craig G.
REGISTRATION NUMBER: 39,044
REFERENCE/DOCKET NUMBER: P1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1489
                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:

APPLICATION NUMBER: 60/027070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 mb floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                            583 LLQNPGQELSFS 594
                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 30-Sep-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/08/953,823A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: South San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET:
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                                                                                                                                                                                                                                                                                        LENGTH:
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                                                                                                                                        1 LLORPGLOLYFS 12
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                                                                                                                                                                                                                                                               OPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 94080
                                                                                                                                                                                                                                                                        1: 793 amino acids
Amino Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            California
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WinPatin (Genentech)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Genentech, Inc
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Pred. No. 41;
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Pred. No.
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5304637-7 Patent No. 5304637

APPLICANT: DORSSERS, LAMBERTUS C.J.; WAGEMAKER, GERARD; VOS.; VONNE J.; VAN LEEN, ROBERT W.; PERSOON, MARIA L.N.
TITLE OF INVENTION: EXPRESSION AND PURIFICATION OF HUMAN

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INTERLEUKIN-3 AND MUTEINS THEREOF
NUMBER OF SEQUENCES: 30
CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/ FILING DATE: 13-MAR-1990

US/07/494,182

NUMBER: 249,184

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Query Match:
Best Local Similarity
-has 7; Conserv:
                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                         REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-
TELECOMMUNICATION INFORMATION
TELEPHONE: (415) 781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/216,260
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
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227 LLRSPGRYVYFS 238
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 08/168,643 FILING DATE: 16-DEC-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 08/030,073 FILING DATE: 19-MAR-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: FILING DATE: 06-DE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: United States ZIP: 94111-4187
                                                                                                                                                                                     STRANDEDNESS:
                                                                                                                                                                                                                                                                                          TELEFAX:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 4
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                              1 LLQRPGLQLYFS 12
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                                                                 Conservative
                                                                                                                                                                                                                                                                                            (415)
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                                                                                                                                                                   linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Flehr, Hohbach, Test, Albritton & Herbert Attn: Walter H. Dreger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-DEC-1993
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                                                                 Mismatches
                                                                                Score 36; DB 2;
Pred. No. 1e+02;
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Best Local Similarity
~+~hes 7; Conserv
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5304637-7
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5304637-1
;Patent No. 5304637
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: DORSSERS, LAMBERTUS C:J.; WAGEMAKER, GERARD; VO. VONNE J.; VAN LEEN, ROBERT W.; PERSOON, MARIA L.N. TITLE OF INVENTION: EXPRESSION AND PURIFICATION OF HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; FILING DATE: SEQ ID NO:7:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO:1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INTERLEUKIN-3 AND MUTEINS THEREOF NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 12, Application US/08284393B
APPLICATION UMBER: US/08/284,393:
FILING DATE: 01-AUG-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Ching, Edwin P.
REGISTRATION NUMBER: 34,090
REFERENCE/DOCKET NUMBER: DX0389
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                    ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                       tent No. 569623
                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 249,184
FILING DATE: 16-AUG-1988
                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Zurawski, Sandra M.
APPLICANT: Zurawski, Gerard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                ORRESPONDENCE ADDRESS:
ADDRESSEE: DNAX Research Institute
                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Zurawski, Gerard
TITLE OF INVENTION: MUTEINS OF MAMMALIAN CYTOKINES
NUMBER OF SEQUENCES: 16
                                                                                                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                      CITY: Palo Alto
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                                                                                                                                                                                                                                                                                                 901 California Avenue
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                                                                                                                                                                                                                                                       USA
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87.5%;
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87.5%;
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Pred. No. 56;
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Pred. No.
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22;
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12 LLVRPGLQ 19

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; MOLECULE TYPE:
US-08-284-393B-12
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PCT-US95-08950-12
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Best Local Similarity
7; Conserva
                                                                                         PCT-US95-08950-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 12, Application PC/TUS9508950 GENERAL INFORMATION:
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TELEFAX: 415-496-1200
INFORMATION FOR SEQ ID NO:
                                                                                                        MOLECULE TYPE:
                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 152 amino acids
                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION: TELEPHONE: 415-852-9196
                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: Ching, Edwin P.
REGISTRATION NUMBER: 34,
                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Zurawski, APPLICANT: Zurawski,
                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: U
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS
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                                                                                                                                                      TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                     TOPOLOGY:
                                                                                                                                    STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                               FILING DATE: 01-AUG-199
                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: DX0389
                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Palo Alto
STATE: California
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                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
STRANDEDNESS: si
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH:
                                                                                                                                                    amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               901 California Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      152 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            linear
                                                                                                                       linear
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                                                                                                                                    single
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87.5%;
                        54.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MUTEINS OF MAMMALIAN CYTOKINES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sandra M.
                                                                                                                                                                                                                                                                                                                                            US 08/284,393
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gerard
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                                                                                                                                                                                                                                                                             34,090
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         Score 33; DB 5;
Pred. No. 57;
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 33; DB
Pred. No. 57;
                                        Length 152;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 152;
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RESULT 11 US-07-952-853-22

Sequence 22, APF-

Application US/07952853

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TOPOLOGY: 1:

MOLECULE TYPE:
US-07-952-853-22
                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
                                                                                                                     Sequence 22,
                                                                                                                                -08-914-848-22
                                                                                                                                                                                                                                                                                                                                                                                                                      NFORMATION FOR SEQ ID NO:
                                                                                                         tent No.
                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS
LENGTH: 628 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 24
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-813-5600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
                                                           APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: Murashige, Kate H.
REGISTRATION NUMBER: 29,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Palo Alto
STATE: California
ZIP: 94304-1018
                                                                                                                                                                                             62 MLQNPGLQ 69
                                                                                                                                                                                                                                                                                                                                                                            TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US
FILING DATE: 19921125
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
                                                                                                                                                                                                               1 LLQRPGLQ 8
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                                                                                       INFORMATION:
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                                                                                                                     Application US/08914848
                                                                                                                                                                                                                                                                                                                                                                                          628 amino acids
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                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                     415-494-0792
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           De Graaff, Leendert H.
                                                        Bakhuis,
                                                                         Van Heuvel, Margaretha
                                                                                                                                                                                                                                                                                                                                                             linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MORRISON & FOERSTER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Floppy disk
                                                                                                                                                                                                                                                                                                                                              protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Jacob
                          Abraham
                                                                                                                                                                                                                                                                      54.18;
75.08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Michel J. A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORIGIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ENCODING ARABINAN-DEGRADING ENZYMES OF FUNGAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Janna G.
Michel J. A.
                                                          Janna G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Peter M.
              Leendert H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US/07/952,853
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                                                                                                                                                                                                                                                                    Score 33; I
Pred. No. 2
                                                                                                                                                                                                                                                       Mismatches
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                                                                                                                                                                                                                                                                                    DB 2;
                                                                                                                                                                                                                                                                      6e+02
                                                                                                                                                                                                                                                                                 Length 628;
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US-08-914-848-22
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Best Local Similarity
                                                                                                                                                                                                                                                                                               Sequence 3, Patent No. (
                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATA:
CLASSIFICATION:
PRIOR APPLICATION NUMBER: US 07.
APPLICATION NUMBER: 25-NOV-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: California
ZIP: 94304-1018
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 628 amino acid
TYPE: amino acid
                                                                                                                                                                                                 TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
                                   COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY DISK
                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION: TELEPHONE: 415-813-5600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 25-NOV-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                 NUMBER OF SEQUENCES:
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS MOS
                                                                                                                                                                                                                                                                                                                                                                                           62 MLQNPGLQ 69
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SOFTWARE: PatentI
                                                                                                    CITY: NEW YORK
STATE: NEW YORK
                                                                                                                                     STREET:
                                                                                   COUNTRY:
                                                                                                                                                 ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                           1 LLQRPGLQ 8
                                                                      10154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Murashige, Kate
                                                                                                                                                                                                                                                                                                            Application US/08840316
                                                                                                                                     345 PARK AVENUE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     628 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       415-494-0792
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                     USA
                                                                                                                                                                                                                                                               Emerson,
                                                                                                                                                    MORGAN & FINNEGAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               IBM PC compatible SYSTEM: PC-DOS/MS-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MORRISON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein
                                                                                                                                                                               Use In Diagnostic Methods And Vaccines
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          54.18;
75.08;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLONING AND EXPRESSION OF DNA
                                                                                                                                                                                                                                                               Suzanne U., Purcell, Robert H.;
     PC-DOS/MS-DOS
                                                                                                                                                                                                                Recombinant Proteins Of A Pakistani Strain Of Hepatitis E And Their
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Peter M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29,959
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22:
                                                                                                                                                                                                                                           A., and Robinson, Robin A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       246152003500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 33; DB 2;
Pred. No. 2.6e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 628
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
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Query Match
Best Local Similarity bu...
Thehes 6; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 758-4800
TELEPAL: (212) 751-6849
NFORMATION FOR SEQ ID NO: 3:
                                                          APPLICATION NUMBER: US 07/681,078
FILING DATE: 05-APR-1991
PRIOR APPLICATION DATA:
                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/279,823
FILING DATE: 25-UUL-1994
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                           SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                          ORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
                                                                                                                                                                            APPLICATION NUMBER: US/08/478,507 FILING DATE: 07-JUN-1995
                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                              TLE OF INVENTION:
 APPLICATION NUMBER:
                             APPLICATION NUMBER: US 07/505,888 FILING DATE: 05 APR-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                            TLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        83 RPGLDLVFA 91
                                                                                                                                                                                                                                                                                                                ZIP: '94306
                                                                                                                                                                                                                                                                                                                            COUNTRY:
                                                                                                                                                                                                                                                                                                                                                          CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                           STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Richard W. Bork REGISTRATION NUMBER: 36 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            , Application US/08478507 6120988
                                                                                                                                                                                                                                                                                                                                                CA
                                                                                                                                                                                                                                                                                                                                                                          3: Dehlinger & Associates
350 Cambridge Avenue, Suite 250
                                                                                                                                                                                                                                                                                                                               USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  123 amino acid residues
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Krawczynski, Krzysztof Z
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Bradley, Daniel W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Reyes, Gregory R
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                                                                                                                                                                                                                                                                            Floppy disk
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US 07/420,921
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US-08-809-523-3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 3, Applica
Patent No. 6207416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO:
                                            APPLICATION NUMBER: US08/316,765 FILING DATE: 03-OCT-1994 PRIOR APPLICATION DATA:
                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/13102
FILING DATE: 03-OCT-1995
                                                                                             FILING DATE: 03-OCT-PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY
                                                                                                                                                                                               CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
TTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                          UMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 123 amino
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
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PRIOR APPLICATION DATA:
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               APPLICATION NUMBER: 07/9 FILING DATE: 18-SEP-1992
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                                                                                                                                                                                                               SOFTWARE:
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ALIGNMENTS

Fc gamma (IgG) receptor high affinity - mouse
C;Species: Mus musculus (house mouse)
C;Date: 02-Jul-1996 #sequence_revision 02-Jul-1996 #text_change 21-Jan-2000

A;Title: Linkage on chromosome 3 of autoimmune diabetes and defective Fc receptor for A;Reference number: I48471; MUID:93242399
A;Accession: I48471

A;Status: preliminary; translated from GB/EMBL/DDBJ

R; Prins, J.B.; Todd, Science 260, 695-698,

Todd, J.A.; Rodrigues, N.R.;

Ghosh, S.;

Hogarth, P.M.; Wicker,

C;Accession: I48471

I4847]

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R;Allen, J.M.; Seed, B. 11824, 1988
Nucleic Acids Res. 16, 11824, 1988
A;Title: Nucleotide sequence of three cDNAs for the human high affinity Fc receptor (A;Reference number: S03018; MUID:89098339
                                                                                                                                                                                                                                                                                                                                                               Fc gamma (IgG) receptor I (high affinity) form b - human
N;Alternate names: CD64
C;Species: Homo sapiens (man)
C;Date: 03-Apr-1992 #sequence_revision 03-Apr-1992 #text_change 21-Jan-2000.
C;Accession: A41357; S03019
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F;128-180/Domain: immunoglobulin homology <IMM>
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                                                                       A; Accession:
                                                                                                                                                                                                       A; Molecule type: mRNA
A; Residues: 1-344 <ALL1>
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Science 243, 378-381,
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A; Residues: 1-336 < RES>
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references: EMBL:X14355; NID:g31333; PIDN:CAA32536.1; PID:g31334
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Seed, B.
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A; Note: the authors translated the codon ACT for re
C; Superfamily: Fc gamma receptor I; immunoglobulin
C; Keywords: alternative splicing; immunoglobulin re
                  C;Keywords: alternative splicing; glycor; 1-15/Domain: signal sequence #status
                                                                                                            A;Cross-references: GDB:135911; OMIM:146760
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                                                                                                                                                                                                                                                                                                                      ;Molecule type: mRNA;Residues: 1-24,'5',26-337,'T',339-374 <AL2>;Residues: 1-24,'5',26-337,'T',339-374 <AL2>;Cross-references: EMBL:X14356; NID:g31331; PIDN:CAA32537.1; PID:g31332;Perez, C.; Wietzerbin, J.; Benech, P.D.
ol. Cell. Biol. 13, 2182-2192, 1993
ol. Cell. Biol. 13, 2182-2192, 1993
;Title: Two cis-DNA elements involved in myeloid-cell-specific expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ;Date: 30-Dec-1991 #sequence_revision 06-Sep-1996 #text_change ;Accession: A39878; I70304; B41357; $03018; I57525 ;Van de Winkel, J.G.J.; Ernst, L.K.; Anderson, C.L.; Chiu, I.M. Biol. Chem. 266, 13449-13455, 1991
                                                              Superfamily: Fc gamma receptor I; immunoglobulin homology
                                                                                                                                                                          Cross-references: GB:S57204; NID:g298692;
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Residues: 1-8 <RE2>
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Cross-references: GB:X14356; GB:M21091; NID:g31331;
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Reference number: A41357; MUID:89100284
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Seed, B.
                                                                                       1q21-1q21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             90, 2102-2109, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16, 11824,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        I55577;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           I-A (high affinity) precursor
                                                                                                                                                                                                                                             GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    eptor I family gene
MUID:93055454
                                                                                                                                                                                                                                                                                        MUID: 93204964
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NID: g184840;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
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Pred. No. 0.00098;
); Mismatches 0;
                                         ycoprotein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CDNAS
predicted <EXT>
                                                                                                                                                                            PIDN: AAD13887.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PIDN:AAA36049.1;
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bulin homology
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                                       immunoglobulin
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                       <SIG>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 344;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PIDN:CAA32537.1;
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                                                                                                                                                                          PID:g4261587
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PID:g292169
                                       receptor;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           mononuclear cells
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                                                                                                                                                                                                                                                                                                                                  in
                  A; Reference number: A; Accession: S06628
  A; Molecule type: DNA
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R;Krasney, P.A.; Carr, C.; Cavener, D.R.

No. Biol. Evol. 7, 155-177, 1990

A;Title: Evolution of the glucose dehydrogenase

A;Reference number: A39019; MUID:90205602

A;Accession: A39019
                                                                                                                                                                                                                                     glucose dehydrogenase (acceptor) (EC 1.1.99.10) - 1
C;Specias: Drosophila melanogaster
C;Date: 27-Nov-1991 #sequence_revision 09-May-1997
C;Accession: A39019; S06628
                                                                                                                                                                                                                                                                                                                                                                                                     В
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C;Species: Mus musculus (house mouse)
C;Date: 18-Jun-1993 #sequence_revision 18-Nov-1994 #text_change 21-Jan-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               F;117-170/Domain: immunoglobulin homology <IMM2> F;293-313/Domain: transmembrane #status predicte F;59,78,152,159,163,195,240/Binding site: carboh
                                                                                                                                                                                                                                                                                                                                                                                                                                         QУ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Cross-references: GB:M31314; NID:9200752; PIDN:AAA40056.1; C;Superfamily: Fc gamma receptor I; immunoglobulin homology C;Keywords: immunoglobulin receptor; transmembrane protein F;127-179/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Title: Structure and mapping of the gene encoding mouse A;Reference number: A46480; MUID:92166399 A;Accession: A46480
                A; Title: Molecular structure and
                                     Genetics 120, 475-484, 1988
                                                      R; Whetten, R.;
                                                        A;Cross-references: GB:M29298; NID:g157499; R;Whetten, R.; Organ, E.; Krasney, P.; Cox-
                                                                                                                                                                                                                                                                                                                            A39019
                                                                                                                                                                                                                                                                                                                                             RESULT
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                                                                                              A; Residues: 1-612 < KRA>
                                                                                                                 A; Molecule type:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Title: Molecular cloning and expression A; Reference number: A43511; MUID: 90111035
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Note: sequence extracted from R; Sears, D.W.; Osman, N.; Tate, J. Immunol. 144, 371-378, 1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R;Osman, N.; Kozak, C.A.; Mo
J. Immunol. 148, 1570-1575;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C; Accession: A46480;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Fc gamma (IgG) receptor high affinity - mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A46480
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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nes 12; Conserv
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ilarity 100.0%;
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91.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            McKenzie, I.F.; Hogarth, P.M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1992
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NCBI backbone (NCBIN:85205, NCBIN:85208, B.; McKenzle, I.F.C.; Hogarth, P.M.
transformation of the :89065357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 56; DB
Pred. No. 0.00
1; Mismatches,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 61;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      site: carbohydrate (Asn) (covalent) #status predi
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                                                        7499; PIDN:AAA28571.1; Cox-Foster, D.; Caven
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0.0097;
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                                                                                                                                                                                                                                                                                                    fruit fly (Drosophila melanogaster)
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                  glucose dehydrogenase
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                                                           PID:g157500
er, D.
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g syndromes.
F;1-32/Domain: signal sequence #status predicted <SIG>
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                                                                                                                                                                                                                                                                                                                                                                   В
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A;Cross-references:
                                                             A; Experimental source: ADTC5
                                                                         A; Molecule type: protein
A; Residues: 528-533;539-545;600-605 <AK2>
                                                                                                          A; Accession: PC4476
                                                                                                                           A; Molecule type: mRNA
A; Residues: 1-793 < AKI>
                                                                                                                                                     A; Accession: JC5539
                                                                                                                                                                                                                                                                                    Smoothened protein precursor - mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Gene: PAB1688
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          submitted to the EMBL Data Library, July 1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          hypothetical protein PAB1688 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Introns:
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                                                                                                                                                                                                                                  ;Species: Mus musculus (house mouse);Date: 02-Sep-1997 #sequence_revision;Accession: JC5539; PC4476
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             anonymous,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Experimental source: strain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cross-references: GB:AJ248286; GB:AL096836; NID:g5458366; PIDN:CAB49934.1; PID:g54584
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Molecule type: DNA
Residues: 1-393 <KAW>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Accession: A75079
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Reference number:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Date: 20-Aug-1999 #sequence_revision 20-Aug-1999 #text_change 28-Jul-2000
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                                            Comment: This protein is used in the conserved targets in Hedgehog signalling
                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Species: Pyrococcus abyssi
                                                                                                                                                                                                                                                                                                                                                                   189
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
7; Conserv
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5; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 45.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 38;
Pred. No.
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A;Title: Nuclectide sequence of the 3' half of A;Reference number: S30236; MUID:93219099
C; Keywords: DNA
F; 19-49/Region:
                                                                                                             A;Cross-references: GB:AE000192; GB:U00096; NID:g1787125; PIDN:AAC73986.1; PID:g17871A;Experimental source: strain K-12, substrain MG1655
                                                                                                                                                                                                                        A;Title: The complete genome sequence of Escherichia coli K-12.
A;Reference number: A64720; MUID:97426617
A;Accession: C64829
                                                                                                                                                                                                                                                                                                                     R;Blattner, F.R.; Plunkett III, G.; Bloch, C.A.; A.; Rose, D.J.; Mau, B.; Shao, Y.
                                                                                                                                                                                                                                                                                                                                                                                     C;Species: Escherichia coli
C;Date: 12-Sep-1997 #sequence_revision 17-Sep-1997 #text_change 29-Sep-1999
                                                                                                                                                          A; Residues:
                                                                                                                                                                           A; Molecule type:
                                                                                                                                                                                                                                                                                                Science 277, 1453-1462, 1997
                                                                                                                                                                                                                                                                                                                                                                C; Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                transcription regulator year - Escherichia coli
                                                                                                                                                                                                                                                                                                                                                                                                                                                             C64829
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                F;125/Binding site: phosphoryl-RNA (Tyr) (covalent) #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             F;495-1010/Product: RNA-directed RNA polymerase #status predicted <POL>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     F;62-494/Product: nuclear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Note:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Molecule type: mRNA
A; Residues: 1-1016 < WUM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C;Date: 19-Mar-J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Variety:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C; Species: zucchini yellow mosaic virus, ZYMV
                                                                                                                                                                                                     A; Status: nucleic acid sequence not shown; translation not shown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C; Accession: S30236
                                                                  A; Gene:
Superfamily: hypothetical protein b1328; Superfamily: hypothetical protein b1328; Keywords: DNA binding; transcription re; 19-49/Region: helix-turn-helix motif
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Note: the nucleotide sequence was submitted to the EMBL Data Library, Superfamily: tobacco etch virus genome polyprotein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Keywords: coat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cross-references: EMBL:X68509; NID:g288233;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Status: nucleic acid sequence not shown; translation not shown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Accession: S30236
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Best Local
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8; Conserv
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                                                                                                                                                                                  DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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inclusion protein a #status predicted <NIA>
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Pred. No.
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Pred. No. 51;
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                                                                                                                                                                                                                                                                                                                                           Perna, N.T.; Burland, V.; Riley,
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G-protein signaling regulator - fission yeast (Schizosaccharomyces pombe) C;Species: Schizosaccharomyces pombe C;Date: 16-May-1996 *sequence_revision 13-Mar-1997 *text_change 21-Jan-2000 C;Accession: T38179; $62427 C;Accession: T38179; $62427 R;Lye, G.; Churcher, C.M.; Barrell, B.G.; Rajandream, M.A.; Walsh, S.V.
                                                                                                                                                        RESULT
S62427
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    hypothetical protein F45C12.14 - Caenorhabditis elegans C; Species: Caenorhabditis elegans C; Date: 29-Oct-1999 #sequence_revision 29-Oct-1999 #tex
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A;Accession: H83705
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; Johnson, D.; Kramer, J.; Keppler, D.
ubmitted to the EMBL Data Library, July 1997
ubeription: The sequence of C. elegans cosmid F45C12.
; Reference number: Z21123
                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local
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Date: 01-Dec-2000 #sequence_revision
                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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72.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 37; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29-Oct-1999 #text_change
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25;
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S50277
                                                                                                                                                  hypothetical protein PA3064 [imported] - Pseudomonas aeruginosa (strain PA01) C;Species: Pseudomonas aeruginosa C;Date: 15-Sep-2000 #sequence_revision 15-Sep-2000 #text_change 31-Dec-2000 C;Accession: G83264
                                                                                                                     C; Accession: G83264
R; Stover, C.K.; Pham, X.Q.;
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                               Nature 406,
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Olson, M.V

Y.; Brody,

Erwin, L.L.; (

Coulter,

Mizoguchi, r, S.N.; Fo

hi, S.D.; Warrener, Folger, K.R.; Kas,

Hickey, Larbig,

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959-964,

genome

sequence of Pseudomonas aeruginosa PA01, an opportunistic

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A;Map position: 5R
C;Superfamily: deubiquinating enzyme
C;Keywords: hydrolase
                                                                                                                                                                                                                                                                                                                  submitted to the EMBL Data Library, A; Description: The sequence of S. cc A; Reference number: S50647
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            N; Alternate names: protein YEK144c
C; Species: Saccharomyces cerevisiae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Experimental source: C; Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Reference number: A; Accession: T38179
                                                                                                                                                                                                  A; Cross-references:
                                                                                                                                                                                                                                                                  A; Molecule type: DNA
A; Residués: 1-805 <DIE>
                                                                                                                                                                                                                                                                                                                                                                    R;Dietrich,
                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Title: UBP5 encodes a putative yeast ubiquitin-specific A;Reference number: S50277; MUID:95176708 A;Accession: S50277
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C; Accession: S50277; S50647
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C;Date: 01-Aug-1995 #sequence_revision 01-Sep-1995 #text_change 29-Sep-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ubiquitin-specific proteinase UBP5 (EC N; Alternate names: protein YER144c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Gene: rgs1; SPAC22F3.12c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Cross-references:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Status: preliminary; translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          submitted to the
                                                                                                                                                                                                                                                                                                    A; Accession: S5064
                                                                                                                                                                                                                                                                                                                                                                                    A;Cross-references: EMBL:U10082;
                                                                                                                                                                                                                                                                                                                                                                                                       A;Residues:
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                                                                                                                                                                                                                     A;Gene: SGD:UBP5
                                                                                                                                                                                                                                                    A;Cross-references:
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Best Local Similarity
Matches 6; Conserv
                                                                                 Query Match
Best Local
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                                                                   Matches
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                                2 LQRPGLQLYFS 12
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VQRPQLPLYYS 347
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                                                                Similarity 7; Conserv
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er: Z21776
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                                                                                                                                                                                                  SGD:S0000946; MIPS:YER144c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EMBL: Z54285; NID: ce: strain 972h-;
                                                                                                                                                                                                                                                EMBL:U18917; NID:g603377; PIDN:AAB64671.1; PID:g603384; MIPS:YER1
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54.5%;
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                                                                                                                                                                                                                                                                                                                                                                                    NID:g595376; PIDN:AAC48928.1; PID:g595377
                                                              Score 37; DB Pred. No. 63; 2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NID:g1008429; PIDN:CAA91077.1; PID:g1008440; GSPDB:G
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Pred. No.
                                                                                                                                                                                                                                                                                                                                   // December
cerevisiae
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A;Accession: G83264
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v;Contains: capsid p
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Matches 6; Conserv
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;Status: preliminary
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Title: Cloning full-length dengue type 4 viral DNA sequences:
Reference number: A94352; MUID:87044106
Accession: A94352
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                                                        Query Match
Best Local Similarity
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MacKow, E.; Makino, Y.; Zhao, B.; Zhang, Y.M.; Markoff, L.; Buckler-White, A.; Guiler, rology 159, 217-228, 1987
Title: The nucleotide sequence of dengue type 4 virus: analysis of genes coding for no Reference number: A94364; MUID:87293881
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Experimental source: strain PAO1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Zhao, B.; Mackow, E.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Genetics:
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Residues: 1-776 <ZHA>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ross-references: GB:M17255
uperfamily: yellow fever virus
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186/Product: nonstructural protein NS4b #status predicted <N4B>
186/Product: nonstructural protein NS4b #status predicted <N4S>
186/Product: nonstructural protein NS5 #status predicted <NS5>
186/Product: nonstructural protein NS5 #status predicted <NS5>
187/Product: nonstructural protein NS4b #status predicted <NS5>
187/Product: nonstructural protein NS5b #status predicted <NS5>
187/Production NS5b #status predict
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43/Product: .nonstructural protein NS2a #status predicted <N2A>
73/Product: nonstructural protein NS2b #status predicted <N2B>
91/Product: nonstructural protein NS3 #status predicted <NS3>
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                                                        60.7%;
75.0%;
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Score 37; DB 1; repred. No. 2.4e+02; oranatches 0;
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Qy 5 PGLQLYFS 12
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Db 466 PGLELYFN 473
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Search completed: June 4, 2001, 12:15:41
Job time: 274 sec

GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.

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Result
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FCG1_MOUSE
POLG_BCMVN
ALR_HELPY
Y824_DEIRA
ATTY_RAT
                             RR8_EPIVI
RR8_TOBAC
RR8_MAIZE
RR8_ORYSA
KDSB_AQUAE
ALR_HELPJ
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YM37_YEAST
RR8_MUSAC
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ALIGNMENTS

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EMBL; X14356; CAA32537.1; EMBL; X14355; CAA32536.1; PIR; S03018; S03018. PIR; S03019; S03019. PIR; A41357; A41357. PIR; B41357; B41357. MIN; 146760; InterPro; IPR003006; Pfam; PF00047; ig; 3.	This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).	SUBCELLULAR LOCATION: TYPE I MEMBRANE PROTE ALTERNATIVE PRODUCTS: 2 ISOFORMS; A (SHOWN PRODUCED BY ALTERNATIVE SPLICING. TISSUE SPECIFICITY: MONOCYTE/MACROPHAGE-SPE - SIMILARITY: CONTAINS 3 IMMUNOGLOBULIN-LIKE DATABASE: NAME-PROW; NOTE-CD guide CD64 ent WWW-"http://www.ncbi.nlm.nih.gov/prow/cd/cd	UENCE FROM LINE-89100 en J.M., Solation ar olation ar plementary ence 243:3 FUNCTION:	SEQUENCE FROM N.A. SEQUENCE FROM N.A. MEDLINE-89098339; PubMed-2974947; Allen J.M., Seed B.; "Nucleotide sequence of three cDNAs for the human high affinity Fc receptor (FCRI)."; nucleic Acids Res. 16:11824-11824(1988).	(Rel. 40), Last TY IMMUNOGLOBULI (IGG FC RECEPTOI CGRI OR FCGI OR S (Human); Mctazoa; Chordat utheria; Primate 9606;	ILT 1 _HUMAN FCG1_HUMAN STANDARD; PRT; 374 AA. F12314; F12315; 01-OCT-1989 (Rel. 12, Created) 01-OCT-1989 (Rel. 12, Last sequence undate)

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                                                                                  "Structure and mapping of the gene encoding morgamma RI and chromosomal location of the human J. Immunol. 148:1570-1575(1992).
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DOMAIN
AFFINITY RECEPTOR.

- SUBCELLULAR LOCATION: TYPE I MEMBRANE PROTEIN
- TISSUE SPECIFICITY: MACROPHAGE-SPECICIC.
- SIMILARITY: CONTAINS 3 IMMUNOGLOBULIN-LIKE C2
                                                                                                                                      Osman N., Kozak C.A., McKenzie I.F.,
                                                                    -1- FUNCTION: BINDS TO THE
                                                                                                                                                      MEDLINE=92166399;
                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Rodentia;
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2C2AA8103ECF16E6 CRC64;
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Pred. No.
                                                                    REGION
                                                                                                                                                                                                                                                                                                                                        Craniata; Vertebrata; | Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EXTRACELLULAR (POTENTIAL)
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                                                                  OF IMMUNOGLOBULINS
                                                                                                                                      Hogarth P.M.;
                                                                                                                                                                                                                                        the mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GLCNAC.
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                                                                                                     mouse
man Fc
                                                                                                                                                                                                                                      high affinity
                                                                                                                                                                                                                                                                                                                                                                                                                          I PRECURSOR (FC-GAMMA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 374;
                                                                                                 high affinity
gamma RI gene
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; Murinae; Mus
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                                                                    GAMMA. HIGH
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                                                                                                                                                                                                                                      FC
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Best Local S
Matches 11
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15-DEC-1998 (Rel. 37, Last sequence update)
15-DEC-1998 (Rel. 37, Last anotation update)
15-DEC-1998 (Rel. 37, Last anotation update)
15-DEC-1998 (Rel. 37, Last anotation update)
GENOME POLYPROTEIN (CONTAINS: N-TERMINAL PROTEIN (P1); HELPER
COMPONENT PROTEINASE (EC 3.4.22.-) (HC-PRO); PROTEIN P3; 6 KDA PROTEI
1 (6K1); CYTOPLASMIC INCLUSION PROTEIN (C1); 6 KDA PROTEIN 2 (6K2);
1 (6K1); CYTOPLASMIC INCLUSION PROTEIN (C1); 6 KDA PROTEIN 2 (6K2);
1 (6K1); CYTOPLASMIC INCLUSION PROTEIN A (NI-A) (NIA)
1 (EC 3.4.22.-) (49 KDA PROTEINASE) (49 KDA-PRO); NUCLEAR INCLUSION
1 (EC 3.4.22.-) (49 KDA PROTEINASE) (49 KDA-PRO); NUCLEAR INCLUSION
1 (COAT PROTEIN (CP)).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CARBOHYD
CARBOHYD
CARBOHYD
CARBOHYD
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DOMAIN
CARBOHYD
                                   common mosaic virus (NL3 strain).";
Virus Res. 39:13-23(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBL; M31314; AAA40056.1;
PIR; A43511; A43511.
PIR; A46480; A46480.
HSSP; P12319; IALT.
                                                            MEDLINE-96191623; PubMed-8607279; Fang G.W., Allison R.F., Zambolim The complete nucleotide sequence
                                                                                                                                                                                                                                                                                                                                              BCMVN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  entities requires a license agreement (s or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            use by non-profit institutions as long modified and this statement is not removed.
                                                                                                   SEQUENCE FROM N.A.
                                                                                                                              NCBI_TaxID=12196;
                                                                                                                                             Potyvirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TRANSMEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CHAIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pfam; PF00047; ig; 3.
IgG-binding protein; Receptor; Transmembrane; Glycoprotein; Signal;
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InterPro; IPR003006; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  between
                                                                                                                                                       /iruses; ssRNA
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         FUNCTION: HELPER COMPONENT-PROTEINASE TRANSMISSION AND ALSO HAS PROTEOLYTIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ween the Swiss Institute of Bioinf
European Bioinformatics Institute
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CAUTION:
                                                                                                                                                                                                                                                                                                                                                                                                               LLQRPGLQLYFS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                        mosaic virus (strain NL-3 / RNA positive-strand viruses,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25
298
321
46
67
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28
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91.7%;
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IG-LIKE C2-TYPE DOMAIN

IG-LIKE C2-TYPE DOMAIN

IG-LIKE C2-TYPE DOMAIN

IG-LIKE C2-TYPE DOMAIN

N-LINKED (GLCNAC...)

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Pred. No. 0.0042;
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EXTRACELLULAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    POTENTIAL
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                                                             E.M
                                                             i genome
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                                                             Maxwell D.P., Gilbertson enome organization of bean
                                                                                                                                                        Michigan) (BCMV).
no DNA stage; Potyviridae;
         IS REQUIRED FOR ACTIVITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          as its content
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2 (6K2);
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Best Local
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01-CCT-2000 (Rel. 40, I
01-CCT-2000 (Rel. 40, I
CLUCOSE DEHYDROGENASE (
GLUCOSE DEHYDROGENASE (
                                                                                         DHGL_DROME (
P18173; Q9VI87;
01-NOV-1990 (Re)
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                                                                                                                                                                                                                                                                                                   CHAIN
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Pfam; PF00680; RNA_dep_RNA_pol; 1.
PRINTS; PR00966; NIAPOTYPTASE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pfam;
Pfam;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              modified and this statement is not removed entities requires a license agreement (See
                                    GLD OR CG1152
                                                                                                                                                                                                                                                                                                                                     CHAIN
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Pfam; PF00851; Peptidase_C6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBL; U19287;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the European Bioinformatics Institute.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEROPS;
             Eukaryota;
                       Drosophila melanogaster (Fruit fly)
                                                                                                                                                                                                                                                                                                                                                                                                                  ATP-binding.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     POLYPROTEIN WHICH UNDERGOES POSTTRANSLATIONAL PROTEOLYTIC PROCESSING RESULTING IN THE PRODUCTION OF AT LEAST EIGHT
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                                                                                                                                                                         VVARPGLQAYF 1988
                                                                                                                                                                                                                                                                                                                                                                                                                              protein; Polyprotein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PF01577; Poty_P1;
                                                                                                                                                                                                                    7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             VPG IS COVALENTLY LINKED TO THE GENOMIC RY THE VIRAL RNA OF POTYVIRUSES IS EXPRESSED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   non-profit institutions as long
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Transferase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INVOLVED IN REPLICATION
              Metazoa; Arthropoda; Tracheata; Hexapoda;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAB02170.1;
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                                                                                                                 STANDARD;
                                                                                                                                                                                                                                                                           3066
1265
AA; 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             license agreement (See http://www.isb-sib.
                                                                                                                                                                                                                               65.6%;
 Endopterygota; Diptera;
                                                                                                                                                                                                                                                                            350385
                                              Last sequence update)
Last annotation updat
[ACCEPTOR] PRECURSOR
[ACCEPTOR] SHORT PROT
                                                                                           Created)
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Pred.
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6 KDA PROTEIN 1.
                                                                                                                                                                                                                                                                                                            NUCLEAR INCLUSION PROTEIN NUCLEAR INCLUSION PROTEIN
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E358955297FA3F59 CRC64;
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30;
                                              PROTEIN]
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 Brachycera; Muscomorpha;
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Durbin K.J., Evangelista C.C., Ferraz C., Ferriera S., Fleischmann W.
A Fosler C., Gabrielian A.E., Garg N.S., Gelbart W.M., Glasser K.,
A Glödek A., Gong F., Gorrell J.H., Gu Z., Guban P., Harris M.,
A Harris N.L., Harvey D., Heiman T.J., Hernandez J.R., Houck J.,
A Hostin D., Houston K.A., Howland T.J., Wei M.-H., Ibegwam C.,
A Hostin D., Houston K.A., Howland T.J., Wei M.-H., Ibegwam C.,
A Hostin D., Houston K.A., Howland T.J., Wei M.-H., Ibegwam C.,
A Hostin D., Kalush F., Karpen G.H., Ke Z., Kennison J.A., Ketchum K.A.
A Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z.,
A Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z.,
A Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z.,
A Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z.,
A Kimmel B.E., Kodira C.D., Kraft C., McDerson D.,
A Kimmel B.E., Kodira B., McIntosh T.C., McLeod M.P., McPherson D.,
A Kimmel B.E., Mushina N.V., Mobarry C., Morris J., Moshrefi A.,
A Merkulov G., Milshina N.V., Mobarry C., Morris J., Moshrefi A.,
A Menut S.M., Moy M., Murphy B., Murphy L., Muzny D.M., Nelson D.L.,
A Melson D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M.,
A Nelson D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M.,
A Palazzolo M., Pittman G.S., Pan S., Pollard J., Puri V., Reese M.G.,
A Reinert K., Remington K., Sampders R.D.C., Scheeler F., Shen H.,
A Spier E., Spradling A.C., Stapleton M., Strong R., Sun E.,
A Syirskas R., Tector C., Turner R., Venter E., Wang A.H., Wang Z.-Y., Wassamman D.A., Weinstock G.M., Weissenbach J.,
A Weinstock G.M., Weissenbach J.,
A Phong F. M., Zhong W., Zhong G., Zhong Q., Zheng L.,
A Zheng X.H., Zhong G., Zhong Q., Zheng L.,
A Shur B., Shu S., Zhong W., Zhong G., Zhong Q., Zheng L.,
A Zheng X. H., Zhong G., Zhong Q., Zheng H., O
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Sutton G.G., Wortman J.R., Yan
Brandon R.C., Rogers Y.-H.C.,
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Krasney P.A., Carr C.M., Cavener D.R.;
"Evolution of the glucose dehydrogenase
                                                       SELENOCYSTEINE Perlaky S., Mei
                                                                                                                                                                                                                                                                                                         Cavener D.R., Krasney P.;
"Drosophila glucose dehydrogenase and yeast alcohol homologous and share N-terminal homology with other
                                                                                                                                                                                                                                                                                                                                                                                                          SIMILARITY TO YEAST ALCOHOL OXIDASE MEDLINE=91163320; PubMed=2002763;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Whetten R., Organ E., Krasney P., Cox-Fo
"Molecular structure and transformation
gene in Drosophila melanogaster.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "The genome sequence of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ballew R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley
Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Wan K.H.,
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Amanatides P.G., Scherer S.E., Li P.W.,
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COFACTOR: FAD
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287:2185-2195(2000)
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N., Zhong W., Zhou X., Zhu S., Zhu
W., Rubin G.M., Venter J.C.;
of Drosophila melanogaster.";
                                                                                                                                                                                      Cavener D.;
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Smith H.O.,
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Q99835;
30-MAY-2000 (Rel. 3
30-MAY-2000 (Rel. 3
01-OCT-2000 (Rel. 4
                                                                                     TISSUE-Embryonic lung;
MEDLINE-97064168; PubMed-8906787;
Stone D.M., Hynes M., Armanini M.
Scott M.P., Pennica D., Goddard A.
de Sauvage F., Rosenthal A.;
                                                                                                                                                                                                                                                                     HUMAN
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                                                        Sonic hedgehog.";
Nature 384:129-134(1996).
                                                                                                                                                                    Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
Xie J., Murone M., L
Lam C.-W., Hynes M.,
de Sauvage F.J.;
                              SEQUENCE FROM N.A., MEDLINE=98082974; PI
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S06628; S06628
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                   Murone M.,
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                             ., AND VARIANTS BCC LEU-535 AND GLN-562 PubMed=9422511;
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Goddard A., Rosent
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GLUCOSE DEHYDROGENASE [ACCEPTOR] SHORT
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        Zhang C., Bonifas J.M., Epstein E.H. Jr.,
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EMBL; AF114819; AAD17202.1;
EMBL; AF114820; AAD17202.1;
MIM; 601500;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "Activating Smoothened mutations in sporadic basal-cell carcinoma.";
Nature 391:90-92(1998).
-i- FUNCTION: G PROTEIN (COUPLED RECEPTOR THAT PROBABLY ASSOCIATES WITTHE PARCHED PROTEIN (PTCH) TO TRANSDUCE THE HEDGEHOG'S PROTEINS SIGNAL BINDING OF SONIC HEDGEHOG (SHH) TO ITS RECEPTOR PATCHED THOUGHT TO PREVENT NORMAL INHIBITION BY PATCHED OF SMOOTHENED
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InterPro; IPR000539; -.
Pfam; PF01534; Frizzled;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      JOINED
                               1;
                               Score 38; DB Pred. No. 19; 1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                           Transmembrane; Glycoprotein;
                                                                                                                                                                                                                                                               3 POTENTIAL.
CYTOPLASMIC (POTENTIAL)
                                                                                                                                                                                                                                                                            EXTRACELLULAR (POTENTIAL)
3 POTENTIAL.
                                                                                                                                        N-LINKED (GLCNAC...)
N-LINKED (GLCNAC...)
N-LINKED (GLCNAC...)
                                                                                                                                                                           CYTOPLASMIC (POTENTIAL)
                                                                                                                                                                                                                    5 POTENTIAL.
CYTOPLASMIC (POTENTIAL):
                                                                                                                                                                                                                                 EXTRACELLULAR (POTENTIAL)
5 POTENTIAL.
                                                                                                                                                                                                                                                       4 POTENTIAL
                                                                                                                                                                                                                                                                                                CYTOPLASMIC (POTENTIAL).
2 POTENTIAL.
                                                                                                                                                                                                                                                                                                                      EXTRACELLULAR (POTENTIAL)
1 POTENTIAL.
                                                                                                                                                                                                                                                                                                                                                                 oncogene; Disease mutation
                                                                                               /FTId=VAR_007848.

R -> Q (IN BCC; SOMATIC MUTATION)
/FTId=VAR_007849.
                                                                                                                                W -> L (IN BCC; SOMATIC
                                                                                                                                                                                                 EXTRACELLULAR (POTENTIAL).
                                                                                                                                                                                                            6 POTENTIAL
                                                                                                                                                                                                                                                                                                                                            SMOOTHENED HOMOLOG
                                                                                      8B4C459B34D13F83 CRC64;
                                                                                                                                                                                     POTENTIAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEMBRANE PROTEIN.
IN BASAL CELL CARCINOMAS (BCC),
                                       DB
19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     There are no restrictions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           as its content
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                                                   Length 787
                                                                                                                                           (POTENTIAL)
(POTENTIAL)
                                                                                                                               MUTATION)
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                                                                                                                                                                                                                                                                                                                                                                            Signal
                                0;
                               Gaps
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RESULT
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Best Local
                                                        30-MAY-2000 (Rel. 39
30-MAY-2000 (Rel. 39
30-MAY-2000 (Rel. 39
SMOOTHENED HOMOLOG (
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30-MAY-2000
30-MAY-2000
                               HOMS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CARBOHYD
SEQUENCE
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Akiyama H., Shigeno C., Hiraki Y., Shukunami C., Kohno H., Akagi P.
Konishi J., Nakamura T.;
"Cloning of a mouse Smoothened cDNA and expression patterns of
"Cloning of a mouse Smoothened cDNA and expression patterns of
hedgehog signalling molecules during chondrogenesis and cartilage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                  P97698;
30-MAY-2000
                                                                                                                                                                                                                 OWS
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Biochem. Biophys. Res. Commun. 235:142-147(1997).

I- FUNCTION: G PROTEIN-COUPLED RECEPTOR THAT PROBABLY ASSOCIATES WITH
THE PATCHED PROTEIN (PTCH) TO TRANSDUCE THE HEDGEHOG'S PROTEINS
SIGNAL. BINDING OF SONIC HEDGEHOG (SHH) TO ITS RECEPTOR PATCHED IS
THOUGHT TO PREVENT NORMAL INHIBITION BY PATCHED OF SMOOTHENED.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        -!- SUBCELLULAR LOCATION: INTEGRAL MEMBRANE PROTEIN.
-!- SIMILARITY: BELONGS TO FAMILY FZ/SMO OF G-PROTEIN COUPLED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
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Rattus norvegicus (Rat)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CARBOHYD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DOMAIN
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                                                                                                                                                                                                                                                                                                                                                                    583
                                                                                                                                                                                                                                                                                                                                                                                                      1 LLQRPGLQLYFS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RECEPTORS
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                                                                                                                                                                                                                                                                                                                                                                    LLQNPGQELSFS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PR00489; FRIZZLED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     coupled receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         )0 (Rel. 3
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39, Last sequence update)
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G PRECURSOR (SMO).
                                                                                       39,
                                                                                                                                                                                                                                                                                                                                                                    594
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                                                        PRECURSOR (SMO)
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Rodentia;
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                                                                                                                                                      Created)
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   N-LINKED
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EXTRACELLULAR (POTENTIAL).
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2 POTENTIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            N-LINKED
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                                                                                                                                                                                                                 PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LINKED (GLCNAC...) (P-LINKED (GLCNAC...) (P-LINKED (GLCNAC...) (P-LINKED (GLCNAC...) (P-1082767876171167 CRC64;
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LLQRPGLQLYFS

12

Best Query Match Matches

Local

8; Conserv

Conservative

62.3%;

Score 38; Pred. No. 1; Mismatc

DB 19;

1;

Length 793;

Mismatches

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Indels

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CARBOHYD
                                                                                                                                                                                                                                                                                                                                                                                                                     use by non-profit institutions as long modified and this statement is not removed entities requires a license agreement (See
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Scott M.P., Pennica D., (
de Sauvage F., Rosenthal
                                                                                                                                                                                                                                                                                                                                                                                                          or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                            between the Swiss Institute of Bioinf
the European Bioinformatics Institute.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This SWISS-PROT entry is copyright. between the Swiss Institute of Bio
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nature 384:129-134(1996)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sonic hedgehog.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "The tumour-suppressor gene patched encodes a candidate receptor for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota;
Mammalia; I
 SEQUENCE
                                                                                                                                                                                                                  DOMAIN.
                                                                                                                                                                                                                                           DOMAIN
                                                                                                                                                                                                                                                                      SIGNAL
                                                                                                                                                                                                                                                                                  Developmental
                                                                                                                                                                                                                                                                                                                                                                              EMBL; U84402; AAB41789.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=97064168; PubMed=8906787;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TISSUE-Embryo;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
                                                                                                                                                                            TRANSMEM
                                                                                                                                                                                                     TRANSMEM
                                                                                                                                                                                                                                                                                                             PRINTS;
                                                                                                                                                                                                                                                                                                                             Pfam; PF01392; Fz;
                                                                                                                                                                                                                                                                                                                                                       InterPro;
                                          CARBOHYD
                                                      DOMAIN
                                                                    TRANSMEM
                                                                                  DOMAIN
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                                                                                                                                                               DOMAIN
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                                                                                                                                                                                                                                                                                                                                                                 InterPro; IPR000024;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   THOUGHT TO PREVENT NORMAL INHIBITION BY PATCHED OF SMOOTHENED THOUGHT TO PREVENT NORMAL INHIBITION BY PATCHED SIGNAL, BINDING OF SONIC HEDGEHOG (SHH) TO ITS RECEPTOR PATCHED THOUGHT TO PREVENT NORMAL INHIBITION BY PATCHED OF SMOOTHENED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MUSCLE, KIDNEY, AND TESTIS.
SIMILARITY: BELONGS TO FAMILY FZ/SMO OF G-PROTEIN COUPLED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NEURAL TUBE, PRE-SOM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SUBCELLULAR LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RECEPTORS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NEURAL TUBE, PRE-SOMITIC MESODERM AND SOMITES, DEV
BUD, GUT, EYE, TESTES, TARTILAGE, MUSCLE, LUNG, E
THYMUS, TONGUE, JAW, TASTE BUDS, TEETH, AND SKIN.
IN MULTIPLE TISSUES INCLUDING HEART, BRAIN, LIVER,
                                                                                                                                                                                                                                                                                                                                      PF01534; Frizzled;
                                                                                                                                                                                                                                                                                                            PR00489;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eutheria;
                                                                                                                                                                                                                                                                                                  coupled receptor;
                                                                                                                                                                                                                                                                                                                                                      IPR000539;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pennica D., Goddard A.,
                                                                                                                                                                                                                                                                                    protein
                                                                                                                                                                                                                                                                                                             FRIZZLED
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  497
87375
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38
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Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              N: INTEGRAL MEMBRANE PROTEIN.
IN EMBRYO, FOUND IN THE EARLY NEURAL
    MΨ.
                                                                                                                                                                                                                                                                                               Transmembrane; Glycoprotein; Signal;
               N-LINKED
                                        N-LINKED
                                                                                             CYTOPLASMIC (POTENTIAL). 6 POTENTIAL.
                                                                                                                                   4 POTENTIAL.
EXTRACELLULAR (POTENTIAL).
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P18479; Q89
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                                                                                                                                                                                                                                                                                                             use by non-profit institutions as long modified and this statement is not removed
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-!- FUNCTION: CYTOPLASMIC INCLUSION PROTEIN HAS HELICASE ACTIVITY.
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15-DEC-1998 (Rel. 37, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
GENOME POLYPROTEIN (CONTAINS: N-TERMINAL PROTEIN (P1);
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PTM: THE VIRAL RNA OF POTYVLRUSES IS EXPRESSED AS A SINGLE POLYPROTEIN WHICH UNDERGOES POSTFRANSLATIONAL PROTECLYTIC PROCESSING RESULTING IN THE PRODUCTION OF AT LEAST EIGHT
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FUNCTION: NUCLEAR INCLUSION PROTEIN A HAS PROTEOLYTIC
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                                                                              PF00863; Peptidase_C4;
PF00851; Peptidase_C6;
                             PF00767; Poty_
PF00680; RNA_C
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15-JUL-1999 (Rel. 38, Last sequence update)
15-JUL-1999 (Rel. 38, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
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16-JUL-1999                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        POLG_ZYMVS
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SIMILARITY: HC PROTEINASE BELONGS TO PEPTIDASE FAMILY C6.
SIMILARITY: NI-A PROTEINASE BELONGS TO PEPTIDASE FAMILY C6.
SIMILARITY: BELONGS TO THE POTYVIRUSES POLYPROTEIN FAMILY
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                                                                                                  SWISS-PROT entry is copyright. It is produced through a collaboration sen the Swiss Institute of Bioinformatics and the EMBL outstation -
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Best Local
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Q89330;
15-DEC-1998 (Rel.
15-DEC-1998 (Rel.
15-DUL-1999 (Rel.
                                                                                                                                                                                                                                   15-DEC-1998 (Rel. 37, Created)
15-DEC-1998 (Rel. 37, Last sequence update)
15-DEC-1998 (Rel. 38, Last annotation update)
15-JUL-1999 (Rel. 38, Last annotation update)
GENOME POLYPROTEIN (CONTAINS: N.-TERMINAL PROTEIN (P1); HELPER
COMPONENT PROTEINASE (EC 3.4.22.-) (HC-PRO); PROTEIN P3; 6 KDA PROTEIN
1 (6K1); CYTOPLASMIC INCLUSION PROTEIN (CI); 6 KDA PROTEIN 2 (6K2);
GENOME-LINKED PROTEIN (VPG); NUCLEAR INCLUSION PROTEIN A (NI-A) (NIA)
(EC 3.4.22.-) (49 KDA PROTEIN (PFG); NUCLEAR INCLUSION
PROTEIN B (NI-B) (NIB) (RNA-DIRECTED RNA POLYMERASE) (EC 2.7.7.48);
COAT PROTEIN (CP)].
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CHAIN
CHAIN
NP_BIND
SEQUENCE
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Pfam; PF00767; Poty_coat; 1.
Pfam; PF00680; RNA_dep_RNA_pol; 1.
Pfam; PF00680; RNA_dep_RNA_pol; 1.
                                                                                                                                                                                                                                                                                                                                                                                               ZYMVR
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PRINTS; PR00966; NIAPOTYPTASE.
Hydrolase; Transferase; Thiol protease; RNA-directed Hydrolase; Transferase; Covalent protein-RNA link;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pfam;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 1972 IINKPGIQAYF 1982
    +++
                                                                                        "Characterization of the P1 protein yellow mosaic virus.";
J. Gen. Virol. 76:37-45(1995).
                                                     -
                                                                                                                                 Wisler G.C., Purcifull D.
                                                                                                                                             MEDLINE-95146958; PubMed-7844540;
                                                                                                                                                         SEQUENCE FROM N.A.
                                                                                                                                                                                                            Viruses; ssRNA positive-strand viruses, no DNA stage;
                                                                                                                                                                                                                        Zucchini yellow mosaic virus (strain Reunion Island) (ZYMV)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CHAIN
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                                                                                                                                                                                   NCBI_TaxID=117129;
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FUNCTION: HELPER COMPONENT-PROTEINASE IS REQUIRED FOR APHID TRANSMISSION AND ALSO HAS PROTEOLYTIC ACTIVITY.
FUNCTION: CYTOPLASMIC INCLUSION PROTEIN HAS HELICASE ACTIVITY.
MAY BE INVOLVED IN REPLICATION.
FUNCTION: NUCLEAR INCLUSION PROTEIN A HAS PROTEOLYTIC ACTIVITY.
PYM: VPG IS COVALENTLY LINKED TO THE GENOMIC RNA.
PYM: THE VIRAL RNA OF POTYVIRUSES IS EXPRESSED AS A SINGLE
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                                                                                                                ., Hiebert E.;
1 protein and coding region
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6 KDA PROTEIN
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HELPER COMPONENT PROTEINASE.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PROTEIN.
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No.
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protein-RNA linkage;
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                                                                                                                    of the zucchini
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Helicase;
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RESULT 11
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Best Local Similarity
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SEQUENCE
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InterPro; IPR00:
InterPro; IPR00:
Pfam; PF00863; I
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Pfam;
SEQUENCE FROM N.A.
STRAIN-K12 / MG1655;
MEDLINE-97426617; PubMed-9278503;
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30-MAY-2000
                                                                                                                                                                                                                                                         P75836;
01-NOV-1997
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                                                                                      NCBI_TaxID=562;
                                                                                                            Bacteria; P
Escherichia
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRINTS; PR00966; NIAPOTYPTASE.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  the European Bioinformatics Institute.
                                                                                                                                                                                                                                                                                                    YCAN_ECOLI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    POLYPROTEIN WHICH UNDERGOES POSTTRANSLATIONAL PROTEOLYTIC PROCESSING RESULTING IN THE PRODUCTION OF AT LEAST EIGHT
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een the Swiss Institute of Bioi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PF00767; Poty_coat; 1:
PF00680; RNA_dep_RNA_pol;
PF00271; helicase_C; 1.
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                                                                                                                               Proteobacteria;
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(Rel. 39, Last annotation update)
TRANSCRIPTIONAL REGULATOR IN DMSC-PFLA
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Poty_P1; 1.
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                                                                                                                               gamma subdivision;
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HELPER COMPONENT PI
                                                                                                                                                                                                                                                                                                    PRT;
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55E51B455C20C537 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           No.
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TO PEPTIDASE FAMILY C4
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01-NOV-1995 (Rel. 3
01-OCT-2000 (Rel. 4
HYPOTHETICAL 54.4 K
SPAC22F3.12C.
                                                                                                                                                                                                                                                 YA8C_SCHPO
Q09777;
01-NOV-1995
                        Lye G., Churcher C.M., Barrell B.G., Rajandream Submitted (SEP-1995) to the EMBL/GenBank/DDBJ da-!- SIMILARITY: CONTAINS 1 RGS DOMAIN.
 This SWISS-PROT entry is
                                                                                                                        Schizosaccharomyces.
NCBI_TaxID=4896;
                                                                                                                                                    Schizosaccharomycetales;
                                                                                                                                                               Schizosaccharomyces pombe (Fission yeast).
Eukaryota; Fungi; Ascomycota; Schizosaccha
                                                                                                                                                                                                                                                                                           SCHPO
                                                                                    STRAIN=97
                                                                                                SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PROSITE; PS00044; HTH_LYSR_FAMILY; 1.

Hypothetical protein; Transcription regulation; DNA-binding.

DNA_BIND 20 40 H-T-H MOTIF (POTENTIAL).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pfam; PF00126; HTH_1; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            or send an email to license@isb-sib.ch)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      use by non-profit institutions as long modified and this statement is not removed. entities requires a license agreement (See
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This SWISS-PROT entry is copyright. It is produced through a between the Swiss Institute of Bioinformatics and the EMBI the European Bioinformatics Institute. There are no restrict
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "A 718-kb DNA sequence of the Escherichia coli K-12 genome corresponding to the 12.7-28.0 min region on the linkage map. DNA.Res. 3:137-155(1996).
-i- SIMILARITY: BELONGS TO THE LYSR FAMILY OF TRANSCRIPTIONAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EcoGene; EG13698; ycaN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Oshima T., Alba H., Baba T., Fujita K., Hayashi K., Honjo A.,
Ikemoto K., Inada T., Itoh T., Kajihara M., Kanai K., Kashimoto I
Kimura S., Kitagawa M., Makino K., Masuda S., Miki T., Mizobuchi
Mori H., Motomura K., Nakamura Y., Nashimoto H., Nishio Y., Saito
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Osnima T., Aiba H., Baba T., Fujita K.,
Ikemoto K., Inada T., Itoh T., Kajihara
Kimura S., Kitagawa M., Makino K., Masu
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Science 277:1453-1474(1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Blattner F.R., Plunkett G. III, Bloch C.A., Perna N.T., Burland V., Riley M., Collado-Vides J., Glasner J.D., Rode C.K., Mayhew G.F.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           InterPro;
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Horiuchi T.;
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55.6%;
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PROTEIN C22F3.12C IN CHROMOSOME
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copyright.
                                                                                                                                                 Schizosaccharomycetaceae;
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Pred. No.
                                                                                                                                                              Schizosaccharomycetes;
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produced through
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                                                               M. A.,
                                                                                                                                                                                                                                                                                                                                                                                                                           Length 302
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C., Yamamoto
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DEPS_YEAST

UBPS_YEAST

P39944;

T 01-FEB-1995 (Rel. 31, Created)

T 01-FEB-1995 (Rel. 31, Last sequence update)

DT 01-NOV-1995 (Rel. 32, Last annotation update)

DE UBIQUITIN CARBOXYL-TERMINAL HYDROLASE 5 (EC 3.1.2.1

"HTOLESTERASE 5) (UBIQUITIN-SPECIFIC PROCESSING PROCESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT
UBP5_YI
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A Dietrich F.S., Mulligan J.T., Hennessey K.M., Allen E., Araujo Aviles E., Berno A., Brennan T., Carpenter J., Chen E., Cherry A Aviles E., Duncan M., Guzman E., Hartzell G., Hunicke-Smith S., A Chung E., Duncan M., Guzman E., Hartzell G., Lew H., Lin D., A Hyman R., Kayser A., Komp C., Lashkari D., Lew H., Lin D., A Mosedale D., Nakahara K., Namath A., Norgren R., Oefner P., Oh Mosedale D., Nakahara K., Namath A., Norgren R., Oefner P., Oh Mosedale D., Nehl P., Schramm S., Shogren T., Smith A Petel F.X., Roberts D., Sehl P., Schramm S., Shogren T., Smith A Taylor P., Wei Y., Yelton M., Botstein D., Davis R.W.;
Submitted (DEC-1994) to the EMBL/GenBank/DDBJ databases.
C -i- CATALYTIC ACTIVITY: UBIQUITIN C-TERMINAL THIOLESTER + H(2)O UBIQUITIN + A THIOL.
C -i- SIMILARITY: BELONGS TO PEPTIDASE FAMILY C19; ALSO KNOWN AS FAMILY 2 OF UBIQUITIN CARBOXYL-TERMINAL HYDROLASES.
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "UBPS encodes a putative yeast ubiquitin-specific protease that is related to the human Tre-2 oncogene product."; yeast 10:1497-1502(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
MEDLINE=95176708; PubMed=7871889;
Xiao W., Fontanie T., Tang M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Saccharomyces cerevisiae (Baker's yeast)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Fungi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE
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DOMAIN 232 31
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PS50132; RGS; 1.
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481 AA;
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RGS.
; FE80CB5E78E88A9D_CRC64;
                                                                                                                                                                                                                                                                                                                                                                               Hennessey K.M., Allen E., Araujo F., Carpenter J., Chen E., Cherry
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This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its

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"In viro processing of dengue virus science".

"In viro processing of dengue virus science".

"In virol. 63:3345-3352(1989).

-!- FUNCTION: THE SMALL PROTEINS NS2A, NS2B, NS4A AND NS4B
HYDROPHOBIC, SUGGESTING A POSSIBLE MEMBRANE-RELATED FUR
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ACT_SITE
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SEQUENCE
                                                                                                                                                                                                                                                                  """ "The nucleotide sequence of dengue type 4 virus: analysis of genes coding for nonstructural proteins.";

Virology 159:217-228(1987).
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Pfam; PF00442;
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                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE OF 774-3386 FROM N.A. MEDLINE-87293881; PubMed-3039728; Mackow E., Makino Y., Zhao B., Zh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       genes coding for structural Virology 155:77-88(1986).
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PS00973; UCH_2_2; 1.
PS50235; UCH_2_3; 1.
n conjugation; Hydrolase;
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                                                                                                                                   PROTEIN M AND GLYCOPROTEIN E. THE NUCLEOCAPSID IS A COMPLEX OF PROTEIN C AND MRNA.
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MEROPS; S07.002;
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PF01350;
PF01349;
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377
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1 113 CAPSID
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  (GLCNAC. .
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PROTEIN E.
PROTEIN NS1.
PROTEIN NS2A.
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Best Local Similarity 75.
Matches 6; Conservative
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_HAEIN
PSTC_HAEIN
P45191;
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01-NOV-1995 (Rel. 32)
01-NOV-1995 (Rel. 32)
PHOSPHATE TRANSPORT (
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STRAIN-RD / KW20 / ATCC 51907;

MEDLINE-95350630; PubMed-7542800;

Fleischmann R.D., Adams M.D., White O., Clayton R.A., Kirkness E.F.

Kerlavage A.R., Bult C.J., Tomb J.-F., Dougherty B.A., Merrick J.M.

MCKenney K., Sutton G., Fitzhugh W., Fields C.A., Gocayne J.D.,

Scott J.D., Shirley R., Liu L.-I., Glodek A., Kelley J.M.,

Weidman J.F., Phillips C.A., Spriggs T., Hedblom E., Cotton M.D.,

Utterback T.R., Hanna M.C., Nguyen D.T., Saudek D.M., Brandon R.C.,

Fine L.D., Fritchman J.L., Fuhrmánn J.L., Geoghagen N.S.M.,

Gnehm C.L., McDonald L.A., Small K.V., Fraser C.M., Smith H.O.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE
PROSITE; PS00402; BPD_TRÂNSP_INN_MEMBR; 1.
Transport; Phosphate transport; Transmembrane; Inner membrane.
DOMAIN 1 22 CYTOPLASMIC (POTENTIAL).
TRANSMEM 23 43 POTENTIAL.
                                                                                                                 modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                       This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Entropean Bioinformatics Institute. There are no restructed by non-profit institutions as long as its content
                                                                                                                                                                                                                                                                         Science 269:496-512(1995).

-i- FUNCTION: PART OF THE BINDING-PROTEIN-DEPENDENT TRANSPORT-
-i- FUNCTION: PROBABLY RESPONSIBLE FOR THE TRANSLOCATION
SUBSTRATE ACROSS THE MEMBRANE (BY SIMILARITY).

-i- SUBCELLULAR LOCATION: INTEGRAL MEMBRANE PROTEIN. INNER MEM
                                                                           EMBL; U32818; AAC23027.1;
TIGR; HI1382; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Haemophilus influenzae.
Bacteria; Proteobacteria;
                                                  Pfam; PF00528; BPD_transp; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NCBI_TaxID=727;
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                                                                InterPro;
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SIMILARITY: WITH INTEGRAL MEMBRANE COMPONENTS OF OTHER BINDING-PROTEIN-DEPENDENT TRANSPORT SYSTEMS. BELONGS TO THE CYSTW
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154 PGLELFFS 161
            5 PGLQLYFS
                         Similarity 75.(6; Conservative
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                         Score 36; DB Pred. No. 18; 2; Mismatches
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Search completed: June 4, 2001, 12:24:04 Job time: 566 sec

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Copyright (c) 1993 - 2000 Comp
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09wy94 drosophila
09w1q7 drosophila
055990 synechocyst
069075 human herpe
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Q05912 zucchini ye
Q9kfn1 bacillus ha
Q16743 caenorhabdi
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Q914t8
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Q94963 drosophila
Q9vi87 drosophila
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ALIGNMENTS

RESULT Q92663 Ş SOR DR RT RAX RPR Query Match Best Local Similarity Matches 12; Conser EMBL; L03418; AAA36049.1; HSSP; P12319; JALT: INTERPRO; IPR003006; PFAM; PF00047; 19; 3. PRODOM; PD002534; -; 1. Q92663; Q92663; 01-FEB-1997 (TrEMBLrel. 01-FEB-1997 (TrEMBLrel. 01-OCT-2000 (TrEMBLrel. SEQUENCE "Novel Fc gamma receptor I family gene products in human mononuclear cells."; MEDLINE-93055454; PubMed-1430234; Porges A.J., Redecha P.B., Doebele R., Kimberly R.P.; TISSUE-BLOOD; Eukaryota; Metazoa; Mammalia; Eutheria; 01-OCT-2000 (TrEMBLrel. FC GAMMA RECEPTOR I. SEQUENCE FROM N.A. NCBI_TaxID=9606 Homo sapiens (Human) Clin. Invest. 90:2102-2109(1992). 1 LLQRPGLQLYFS 12 ilarity 100.0%; Score 61; DB 4; ilarity 100.0%; Pred. No. 0.0013; Conservative 0; Mismatchic 374 AA; PRELIMINARY; Chordata; Craniata; Vertebrata; Euteleostomi; Primates; Catarrhini; Hominidae; Homo. 42632 MW; 15, Created) Last sequence update) Last annotation update) PRT; D33D59398CEEA699 CRC64; 374 Pan L.C., A Length 374; Salmon J.E., Indels ç. Gaps

RESULT 2 Q92495 ID ·Q92495

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              Yan Y., Li X., Wang A., Zhang G.;
"Molecular cloning and identification of full-length cDNA high affinity Fc receptor for bovine IgG (Fc gamma RI).";
Vet. Immunol. Immunopathol. 75:151-159(2000).
EMBL; AF162866; AAF80477.1;
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FC GAMMA RECEPTOR I.
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"Definition of interferon gamma-response elements in a novel human Fc
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01-OCT-2000 (TrEMBLrel. 15, I
FC GAMMA RECEPTOR TYPE I (FC
Receptor.
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PubMed=1402657;
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Catarrhini; Hominidae;
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01-MAY-2000
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                                     Drosophila melanogaster (Fruit fly).
Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;
Eukaryota; Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
NCBI_TaxID=7227;
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Q94963;
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MEDLINE=20196006;
        SEQUENCE FROM N.A. STRAIN-BERKELEY;
                                                                                                      GLD PROTEIN.
                                                                                                                                                                                                                                                                                                          PROSITE; PS00623; GMC_OXRED_1; PROSITE; PS00624; GMC_OXRED_2; PRODOM; PD004359; -; 1.
SEQUENCE 612 AA; 66794 MW;
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EMBL; U63325; AAB48020.1; --
EMBL; U63324; AAB48020.1; JOINED.
EMBL; U63324; AAB48020.1; JOINED.
FLYBASE; FBgn0013875; Dsim\Gld.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A. STRAIN=CAS 54;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Drosophila simulans (Fruit fly).
Eukaryota; Metazoa; Arthropoda; Tracheata; He
Pterygota; Neoptera; Endopterygota; Diptera;
Ephydroidea; Drosophilidae; Drosophila.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GLD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GLUCOSE
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01-OCT-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE-97019051; pubMed=8865667;
Hamblin M.T., Aquadro C.F.;
"High nucleotide sequence variati
in Drosophila simulans is consist
                                                                                                                                                                                                                                                                                                                                                     PFAM; PF00732; GMC_oxred; 1.
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PubMed=10731132;
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77.8%;
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Pred. No. 7.5;
2; Mismatches
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Pred. No.
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Jalali M., Kalush F., Karpen G.H., Ke Z., Kennison J.A., Ketchum K.A.,

Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lal Z.,

Lasko P., Lei Y., Levitsky A.A., Li J., Li Z., Liang Y., Lin X.,

Lasko P., Lei Y., Levitsky A.A., Li J., Li Z., Liang Y., Lin X.,

Lasko P., Lei Y., Levitsky A.A., Li J., Li Z., Liang Y., Lin X.,

Lasko P., Lei Y., Levitsky A.A., Li J., Li Z., Liang Y., Lin X.,

A Liu X., Mattei B., McIntosh T.C., McLeod M.P., McPherson D.,

A Merkulov G., Milshina N.V., Mobarry C., Morris J., Moshrefi A.,

A Mextulov G., Milshina N.V., Mobarry C., Morris J., Moshrefi A.,

A Mount S.M., Moy M., Murphy B., Murphy L., Muzny D.M., Nelson D.L.,

RA Nelson D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M.,

Palazzolo M., Pittman G.S., Pan S., Pollard J., Puri V., Reese M.G.,

RA Reinert K., Remington K., Sunders R.D.C., Scheeler F., Shen H.,

RA Shue B.C., Siden-Kiamos I., Simpson M., Skupski M.P., Smith T.,

RA Spier E., Spradling A.C., Stapleton M., Skupski M.P., Smith T.,

Saylirkas R., Tector C., Turner R., Venter E., Wang A.H., Wang X.,

RA Wang Z.-Y., Wassarman D.A., Weinstock G.M., Weissenbach J.,

RA Wang Z.-Y., Wassarman D.A., Weinstock G.M., Weissenbach J.,

RA Wellams S.M., Moodage T., Worley K.C., Wu D., Yang S., Yao Q.A.,

RA Leng X.H., Zhong F.N., Zhong W., Zhou S., Zhou Q., Zheng L.,

"The genome sequence of Drosophila melanogaster.";

"The genome sequence of Drosophila melanogaster.";
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Best Local Similarity //...
7; Conservative
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Q9UZX2;
01-MAY-2000
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Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
de Pablos B., Delcher A., Deng Z., Mays A.D., Dew I., Dietz S
                                                                                                                                                                                                  Pyrococcus abyssi.
Archaea; Euryarchaeota;
NCBI_TaxID=29292;
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01-JUN-2000
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PROSITE; PS00624; GMC_OXRED_2;
SEQUENCE 612 AA; 66835 MW;
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Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier
Burtis K. D. Brokstein P., Brottier
                                                                                                                                                                                                                                                                                                                                 PAB1688.
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                                                                                                                          SEQUENCE FROM N.A
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Pyrococcus abyssi genome sequence: insights into archaeal
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O (TrEMBLrel. 13,
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C., Blazej R.G., Champe M., Pfeiffer B.D.,
E.G., Helt G., Nelson C.R., Miklos G.L.G.,
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Center A., Chandra
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.A., Galle R.F.
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    chromosome
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Q9KFN1;

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Submitted (JUL-1999) to the EN
EMBL; AJZ48286; CAB49934.1; -
Hypothetical protein.
SEQUENCE 393 AA; 44932 MW;
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Wu M. Yeong C.Y., Lee S.C., Wong S.M.;
"Nucleotide sequence of the 3' half of zucchini yellow mosaic
(Singapore isolate) genome encoding the 4K protein, protease,
                                                                                                                                                   Polyprotein; Covalent protein-RNA NON_TER 1 1
                                                                                                                                                                                                                      PFAM;
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01-OCT-2000
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                                                                                                                         SEQUENCE
                                                                                                                                                                              Hydrolase;
                                                                                                                                                                                            PRINTS; PR00966; NIAPOTYPTASE
                                                                                                                                                                                                                                                                                         -!- SIMILARITY: TO OTHER POTYVIRUSES POLYPROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                               Nucleic Acids Res. 21:1317-1317(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      189
                                                                                                                                                                                                                                    INTERPRO;
                                                                                                                                                                                                                                                                                                                                                                                                                                             polymerase and coat protein."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=117130;
179 IINKPGIQAYF 189
                           1 LLQRPGLQLYF 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 LLQRPGLQLYF
                                                                                                                                                                                                                                                                                                                  PTM: THE VIRAL RNA OF POTYVIRUSES IS EXPRESSED AS A SINGLE POLYPROTEIN WHICH UNDERGOES POSTTRANSLATIONAL PROTEOLYTIC PROCESSING RESULTING IN THE PRODUCTION OF AT LEAST EIGHT
                                                                                                                                                                                                                                                                                                                                                          IT, VPG, IS COVALENTLY LINKED TO PROTEOLYTIC ACTIVITY.
                                                                                                                                                                                                                                                                                                                                                                                        FUNCTION: NUCLEAR INCLUSION PROTEIN A, OR THE N-TERMINAL PART
                                                                                                                                                                                                                                                                                                                                                                                                    MAY BE INVOLVED IN REPLICATION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LMTKPGIEVYF 199
                                                                                                                                                                                                         PF00680; RNA_dep_RNA_pol;
PF00863; Peptidase_C4; 1.
                                                                                                                                                                                                                                                                            X68509; CAA48521.1;
                                                    l Similarity
5; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
5; Conser
                                                                                                                                                                                                                                    IPR001254;
IPR001730;
                                                                                                                                                                                                                                                               IPR001205;
                                                                                                                                                               Protease; RNA-directed RNA polymerase; Coat protein; n; Covalent protein-RNA linkage.
                                                                                                                       1016
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                                                                                                                                    1016
                                                                62.3%;
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01, Last sequence 15, Last annotation
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                                                                   Score 38;
Pred. No.
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nd viruses, no DNA stage; Potyviridae;
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32;
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Length 425

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Best Local
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01-JAN-1998 (TrEMBLrel.
01-JAN-1998 (TrEMBLrel.
01-JUN-2000 (TrEMBLrel.
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ABC TRANSPORTER (
 EMBL; AF016684; AAB66212.1; INTERPRO; IPR001810; -. PFAM; PF00646; F-box; 1. SEQUENCE 425 AA; 48488 M
                                                                                                                                    [2]
SEQUENCE FROM N.A..
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MEDLINE=94150718; PubMed=7906398;
                                                                                                                                                                                                                                                                                                                                                                                                     Caenorhabditis elegans.
Caenorhabditis elegans.
Metazoa; Nematoda;
                                                                         SEQUENCE FROM N.A.
STRAIN-BRISTOL N2;
                                                                                                                                                                                                         Thierry-Mieg J., Thomas K., Vaudin M., Vaughan K., Watton A., Weinstock L., Wilkinson Sproat J., Wohldman
                                                                                                                                                                                                                                                                                                        Wilson R., Ainscough R., Bonfield J., Burton J., C
                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Nemato
Rhabditidae; Peloderinae;
                                                   Submitted
                                                                Waterston
                                                                                                              Submitted (AUG-1997)
                                                                                                                         Johnson D.,
                                                                                                                                                                        Nature 368:32-38(1994).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          016743
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Submitted (MAR-2000) to the
EMBL, AP001508; BAB04167.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bacillus/Staphylococcus
NCBI_TaxID=86665;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATP-binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bacteria; Firmicutes; Bacillus/Clostridium group; Bacillus/Staphylococcus group; Bacillus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRAIN=C-125 / JCM 9153;
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                                                                                                                                                                                       .2 Mb of contiguous nucleotide sequence
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                                                  (JUL-1997) to the
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                                                                                                                         Kramer J.,
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(TrEMBLrel. 15,
(TrEMBLrel. 15,
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                                                                                                           Keppler D.;
the EMBL/GenBank/DDBJ databases
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  MW;
                                                                                                                                                                                                                                                                                                                     Anderson K.,
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Caenorhabditis.
                                               EMBL/GenBank/DDBJ
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Pred. No. 38;
2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Last sequence update)
Last annotation updat
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D42932E3A29F6A7D CRC64;
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RESULT
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Q9K451
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Best Local S
Matches 8
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Best Local
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01-MAY-2000
01-JUN-2000
                                                                                                                                                          Q9VY94;
Q9VY94;
                                                             Drosophila melanogaster (Fruit fly).

Bukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda;

Pterygota; Neoptera; Endopterygota; Diptera; Brachyce

Ephydroidea; Drosophilidae; Drosophila.
                               [1]
SEQUENCE FROM N.A.
                                                                                                         GRIP91.
                                                                                                                    GRIP91
                                                                                                                                                                                                                                                                                                                 "A set of ordered cosmids and a cithe 8 Mb Streptomyces coelicolor Mol. Microbiol. 21:77-96(1996).
EMBL; AL359215; CAB94657.1; -.
SEQUENCE 573 AA; 64041 MW; 1F
MEDLINE-20196006; PubMed-10731132; Adams M.D., Celniker S.E., Holt R.A.,
                       STRAIN-BERKELEY;
                                                     NCBI_TaxID=7227;
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01-0CT-2000 (TIEMBLEEL 15, Last
01-0CT-2000 (TIEMBLEEL 15, Last
CYTOCHROME C OXIDASE POLYPEPTIDE
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Thomson N.R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRAIN=A3(2);
Seeger K.J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Streptomyces coelicolor.
Bacteria; Firmicutes; Actinobacteria; Actinobacteridae;
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01-OCT-2000
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Kinashi H., Hopwood D.A.,
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                                                                                                                   PROTEIN.
                                                                                                                                                                                                                                                               Similarity 7; Conserv
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i (JUN-2000)
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                                                                                                                                                                    PRELIMINARY;
                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                 Parkhill
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e EMBL/GenBank/DDBJ databases.
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Pred. No.
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Pred. No. 53;
0; Mismatches
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                                                                                                                                                                    917
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73;
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chromosome.";
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53;
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                                                                        Brachycera; Muscomorpha;
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Evans

C.A.,

Gocayne J.D.,

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Best Local Similarity bo.,
Best Local Similarity bo.,
Conservative
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Q9W1Q7;
01-MAY-2000
01-MAY-2000
01-JUN-2000
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EMBL, AE003493, AAF48309.1; -
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                                                                                                                                                                    Drosophila melanogaster (Fruit fly)
Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda;
Pterygota; Neoptera; Endopterygota; Diptera; Brachyce
Ephydroidea; Drosophilidae; Drosophila.
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MEDLINE-20196006; PubMed-10731132; Adams M.D., Celniker S.E., Holt R. Amanatides P.G., Scherer S.E., Li
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                                                                             STRAIN-BERKELEY;
                                                                                                 SEQUENCE FROM N.A.
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917 AA; 103706 MW; 6AEE88C211D256BB
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Doyle C., 1
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n J.R., Yandell M
s Y.-H.C., Blazej
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Last annotation updat
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  Evans C.A.,
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Gocayne J.
A., Galle
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Best Local
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Q55990;
Q1-NOV-1996
Q1-NOV-1996
Q1-JUN-2000
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"Sequence analysis of the genome of the unicellular cyanobacterium Synechocystis sp. strain PCC6803. I. Sequence features in the 1 Mb region from map positions 64% to 92% of the genome.";
                                                                MEDLINE-96127529; PubMed-8590279;
Kaneko T., Tanaka A., Sato S., Kotani H.,
Sugiura M., Tabata S.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ballew R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M., Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S.,
                                                                                                                                         SEQUENCE FROM N.A. STRAIN=PCC6803;
                                                                                                                                                                                                                             Synechocystis sp. (strain Bacteria; Cyanobacteria;
                                                                                                                                                                                                                                                                               SLL0743
                                                                                                                                                                                                                                                                                                  HYPOTHETICAL
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64 AA; 7304 MW; 0A
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(TrEMBLrel.
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Pred. No.
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OA31867ED48B244C CRC64;
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                                                                                            Sazuka
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                                                                                          T., Miyajima
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H.O.,
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MEDLINE=97061201; PubMed=8905231; MEDLINE=97061201; PubMed=8905231; Ta

Tanaka A.,

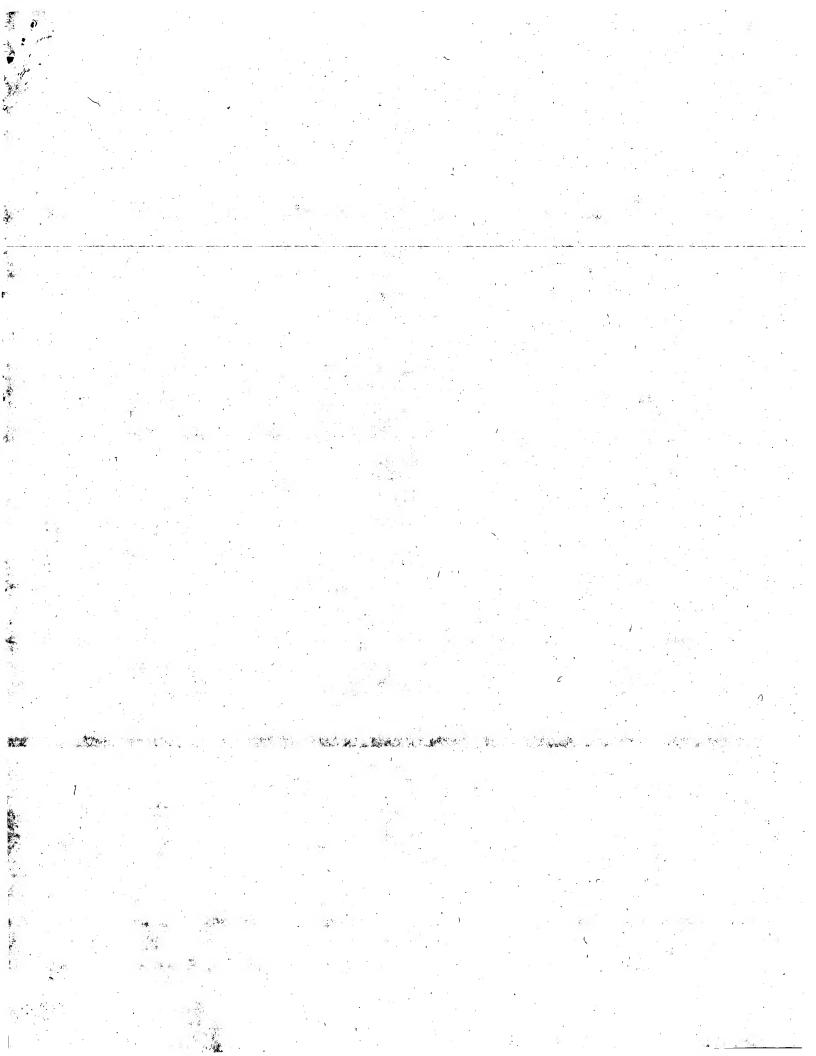
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Best Local Similarity 63.6
...thes 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                     Query Match 59.0%;
Best Local Similarity 58.3%;
Matches 7; Conservative
Q9Y477 PRELIMINARY;
Q9Y477;
01-NOV-1999 (TrEMBLrel. 12, C:
01-NOV-1999 (TrEMBLrel. 12, L:
01-NOV-1999 (TrEMBLrel: 12, L:
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Q69075;
01-NOV-1996 (TrEMBLrel. 01, C
01-NOV-1996 (TrEMBLrel. 01, L
01-NOV-1998 (TREMBLREL. 08, L
GLYCOPROTEIN H (FRAGMENT).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NON_TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "Excretion of non-infectious virus particles lacking glycoprotein H by a temperature-sensitive mutant of herpes simplex virus type 1: evidence that gH is essential for virion infectivity.";

J. Gen. Virol. 69:1147-1156(1988).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          human herpesvirus 1.
Viruses; dsDNA viruses, no RNA stage;
Alphaherpesvirinae; Simplexvirus.
NCBI_TaxID=10298;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
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entire genome and assignment of po
DNA Res. 3:109-136(1996).
EMBL; D64005; BAA10772.1; -
INTERPRO; IPR002636; -
IPFAM; PF01724; DUF29; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=88258458; PubMed=2838568;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRAIN=TSQ26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Desai P.J.,
                                                                                                                                                                                                                                                             123 LLRSPGRYVYFS 134
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hypothetical protein.
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94 LLESPSLQKYF 104
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166 AA;
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17805 MW;
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Pred. No.
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Pred. No. 30;
1; Mismatches
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                                                                                                                      171 AA
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30;
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Kimura T.,
K., Okumura S
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                             Query Match
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Matches 7
                                                             Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                           DJ159A15.1 (FRAGMENT).
DJ159A15.1.
                                                                                                                       NCBI_TaxID=9606;
109 LLKRPQLELY 118
               1 LLQRPGLQLY 10
                              Similarity 70.07; Conservative
                                    59.0%;
70.0%;
                              2
                             Score 36; DB Pred. No. 32; 2; Mismatches
                                   DВ
32;
                                            Length 171;
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Search completed: Job time: 587 sec June 4, 2001, 12:23:24



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Title:
Perfect score:
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Copyright (c) 1993 - 2000 Compugen Ltd.
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R20811
R22550
                                                                                                                                                                    W60556
Y33183
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                                                                                                                                R91439
                                                                                                                                                    R22549
                                                                                                                                                                                                                                                                                    SUMMARIES
                              Human macrophage-s
Human macrophage-s
Human macrophage-s
Human FCRI (CDNA C
Human FCRI (CDNA C
                                                                                                                                                                                                                            Description
                                                                                                                              Human macrophage-s
Human FcRI (cDNA c
                                                                                                                                                                    Human
                                                                                                                                                                                    Oligopeptide from
                                                                                                                                                                      sfc-gammaRI
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56	142	.в21433		G14207	G14208	G14209	R54218	R66462	R38470	в29019	R92705	R44514	G38488	G37164	G28965	G07160	G38489	G37165	G28966	G07161	G38490	G37166	G28967	G07162	в01787	R07302	Y87523	R12428	W86195	в43683	Y96226	961	W97834
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à			Cone snail alpha-	Arabidopsis thalia		Arabidopsis	'L.lactis branched	ALL-1 (acute lymph	ALL-1 protein.	an MLL gene	cDNA clone	MLL amino acid seg		is		thaj	thal	tha	idopsis tha	Arabidopsis thalia	Arabidopsis thalia				Escherichia coli Y	C-terminal sequenc	Conotoxin peptide	Hybrid Fc(gamma)RI	Human Fc receptor	Human cancer assoc	Human high affinit	Human macrophage-	Human Fc receptor

ALIGNMENTS

RESULT W60556 identifying peptide(s) binding specifically to protein target - by expressing on phage surface and testing for binding to immobilised oligo; peptide derived from the target, useful for, e.g. identifying specific antibodies WPI; 1998-240964/21. De Kruif CA, 08-OCT-1996; 07-OCT-1997; WO9815833-A1. Synthetic. immunohistochemical;
cell transformation; Extracellular domain; CD64; identification; antibody; Oligopeptide from extracellular domain of CD64. 18-AUG-1998 W60556; W60556 standard; peptide; 12 (UYUT-) RIJKSUNIV UTRECHT. 16.-APR-1998 (first entry) Logtenberg 96EP-0202791. 97WO-NL00557 immunofluorescent A A analysis; detection;

e.g. identifying

Example

1; Page

29; 40pp; English

of CD64.

Deptides W60537-61 are derived from the They were synthesised on 25 polyethyler

polyethylene

extracellular rods

W60537-61

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Best Local Similarity
watches 12; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 adjacent, non-overlapping sets. Peptide specific phages were isolated by CC affinity binding to the CD64-covered rods. The specification describes a comethod for the identification of a peptide able to bind specifically to a C target protein. The method comprises displaying the peptide on the CS surface of a replicable display package, synthesising oligopeptides concerned from the target protein on a solid phase, and testing for binding between the peptide and oligopeptides. The method is used to screen large peptide libraries, especially to detect antibodies, or their fragments, that bind to cell markers or that can differentiate between different forms of the same protein, including bispecific antibodies that bind to epitopes on the same monomeric antigen or two non-overlapping epitopes on the same monomeric antigen or two cepitopes on different molecules. The genes/oligonucleotides that encode selected peptides can be isolated and used for recombinant production of antibodies. Selected antibodies are used, e.g. for immunohistochemical or immunofluorescent analysis, and also to detect cell transformation caused to monomeric that in a fartivaccanaes
                                                               WPI; 1999-539978/45
                                                                                                 Powell MS;
                                                                                                                Maxwell KF,
                                                                                                                               Baell JB,
                                                                                                                                                                                              11-SEP-1998;
06-FEB-1998;
                                                                                                                                                                                                                                             04-FEB-1999;
                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                         anti-inflammatory;
                                                                                                                                                                                                                                                                                                                                                                                     tissue damage; hypersensitivity; infiniflammatory modulator; Fc-gammaRIa;
                                                                                                                                                                                                                                                                                                                                                                                                                      drug design; therapy; inhibitor;
                                                                                                                                                                                                                                                                                                                                                                                                                                         atomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fc receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human sFc-gammaRI protein fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Y33183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Y33183 standard; Protein;
                                                                                                                                                              (ILEX-) ILEXUS PTY LTD.
                                                                                                                                                                                                                                                                             12-AUG-1999
                                                                                                                                                                                                                                                                                                           WO9940117-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-NOV-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               by mutation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 FYMGSKTLRGRN 12
                                                                                                                                                                                                                                                                                                                                                                                                                               sptor; Fc-gammaRIIa; human; FCR; model; three-dimension; coordinate; bicactive compound design; computer-assisted.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          in (anti)oncogenes.
                                                                                                              McCarthy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                            98US-0099994
98US-0073972
                                                                                                                                                                                                                                             99WO-IB00367
                                                                                                                            Garrett
                                                                                                                                                                                                                                                                                                                                                                         immunoprotective;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Score 64; 100.0%; Pred. No.
                                                                                                              TD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   261 AA.
                                                                                                                               TPJ,
                                                                                                           TPJ, Hogarth PM, Matti
McKenzie IFC, Pietersz
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                   Fc-gammaR; Fc-epsilonR; IgG; IgE; inflammatory cell recruitment; RIa; immune function regulation;
                                                                                                                                                                                                                                                                                                                                                                       sfc-gammaRI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 19;
6.5e-06;
                                                                                                                          Matthews
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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Three-dimensional structures and models of Fc receptors, useful computer-assisted drug design

in

This amino acid sequence was predicted from the cDNA sequence of cDNA clone p98/X2. It differs from the sequence predicted from clone p135 (see Q211/8) at position 25, p135 encodes a Ser residue while clone p98/X2 predicts a Thr residue. At position 58, p90

New CD53 cell surface antigen and DNA encoding it - for immuno-therapy and diagnosis of haematopoietic neoplasms,

etc.

Example 10;

Page 94a; 160pp;

English.

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Matches 12; Conser
                                                                                                                                                                            WPI; 1992-056864/07
                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human macrophage-specific FcRI receptor encoded by clone p98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R22549 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               an FC receptor protein using various computer programs and models. The products of the invention have anti-inflammatory and immunoprotective activity. This sequence represents the human sFC-gammakI protein fragment described in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         hypersensitivity, recruitment of inflammatory cells or release of inflammatory modulators. The therapeutic compositions can also be used to treat disease in animals. The knowledge of the 3-D structure of Fc-gamma RIa and models of other FcR provides a means for designing and producing compounds that regulate immune function and inflammation in an animal, including humans (i.e. structure based drug design). For example, chemical compounds can be designed to block binding of immunoglobulin to
                                                                                                                                                              N-PSDB; Q21179
                                                                                                                                                                                                                                            (GEHO-) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                             13-JUL-1990;
                                                                                                                                                                                                                                                                                                           15-JUL-1990;
                                                                                                                                                                                                                                                                                                                                            23-JAN-1992
                                                                                                                                                                                                                                                                                                                                                                          W09201049-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rapid immunoselection cloning technique; cell surface antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          R22549
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         representing a three-dimensional (3-D) structure that substantially conforms to the specified atomic coordinates. Computer model images of the FCR can be used to design bioactive chemical compounds, e.g. oligonucleotides, peptides, peptides metalics and small organic molecules, oligonucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                        immunodiagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               that inhibit the activity of Fc-gammaR or Fc-epsilonR can be used to reduce IgG- or IgE-mediated tissue damage, e.g. IgG- or IgE-mediated hypersensitivity, recruitment of inflammatory cells or release of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 11; Page 316-317; 326pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   by computer-assisted methods of drug design. Therapeutic compositions
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      261
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                                                                                                                                                                                                                                                                                                           90WO-US04986
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 describes a novel model
                                                                                                                                                                                                                                                                                                                                                                                                                                        high affinity receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        221
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12
                                                                                                                                                                                                              Amiot M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        344
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Pred. No. 0.00019;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              model of an Fc receptor (FcR) protein (3-D) structure that substantially
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 261;
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RESULT
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Best Local
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                  The amino acid sequence (R91439) of human FCRI was detd. from a CDNA clone, p98 (714718), obtd. from a CDNA library using an immunoselection cloning method. FCRI is a high affinity receptor for the FC portion of IgG, normally located on cell surfaces of macrophages. 2 Other cDNA clones, p135 (714717) and p90 (714719),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (see Q21180) predicts Val and p98/X2 predicts Leu. Sequences predicted from all 3 clones show the typical features of a type I integral membrane protein and include a short hydrophobic signal sequence, a single 21 residue hydrophobic membrane spanning domain sequence.
                                                                                                                                                                          Cloning of cDNA encoding cell surface antigen of diagnostic and therapeutic proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cell surface antigen; cloning; immunoselection; therapy; diagnosis; vector; FcRI; Fc receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R91439
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    R91439 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          and a short, highly charged cytoplasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and six Cys residues distributed among three C2 set Ig-related domains A fusion protein of FCRI and a receptor ligand will be helpful to increase the potency of antibodies in therapy.
macrophages. 2 Other cDNA ciones coded for a variants of the FCRI
                                                                                                                                                                                                                                        N-PSDB;
                                                                                                                                                                                                                                                          WPI; 1996-200279/20
                                                                                                                                                                                                                                                                                                Aruffo
                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-DEC-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human FcRI (cDNA clone p98 product)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30-OCT-1996
                                                                                                                                      Example 10; Column 55-56; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                    (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                            18-OCT-1993;
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                                                                                                                                                                                                                                                                                                                                                                                            13-JUL-1989
13-JUL-1990
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                                                                                                                                                                                                                                                                                              Seed B;
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ilarity 100.0%;
Conservative C
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93US-0139273.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "amino acid 58 is Leu in clone p135
translated product"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 64; DB 13;
Pred. No. 0.00025;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
    the C-terminal sequence
                                                                                                                                                                                                useful for isolation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          immunotherapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 344;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
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Y96183
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 100.
Best Local Similarity 100.
Matches 12; Conservative
                                                                                                                                                                                                            /note-
Misc-difference 82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         and p90 products (see also R91438 and W00859). A of FCRI and a receptor ligand will be helpful in potency of antibodies in therapy.
                                                                                                              /note=
Misc-difference 126
                                                                                                                                                                                         /note=
Misc-difference 116
                                                                                                                                                                                                                                                   /note-
Misc-difference 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
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Misc-difference 123
                                                                                                                                                            Misc-difference
                                                                                                                                                                                                                                                                                                    Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                 amyloidosis; multiple sclerosis; parasitic disease; Fc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human macrophage-specific FcRI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Y96183 standard; Protein; 344 AA.
                                                          Misc-difference
                                                                               Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                            panning; immunodiagnosis; diagnosis; immunotherapy; gene th
immune disorder; infection; asthma; immune-complex disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Macrophage; FCRI; cell surface antigen; human; immunoselection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19-DEC-2000
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                                                                                                                                                                                                                                                                                                                     /note=
51
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/note=
213
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134
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117
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50
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Pred. No.
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          AAT"
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                                                                                                                                                                                                                               CAG"
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0.00025;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 344;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    therapy;
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229

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13-JUL-1989;
23-MAR-1990;
13-JUL-1990;
                                                                                                                                                 based upon transient expression of a CSA in eukaryotic
                                                                                                                                                                                                                                            Isolated nucleic acid molecule encoding the CD19 cell surface antigen, useful for immunodiagnosis and immunotherapy of immune-mediated
                                                                                                                                                                                                                                                                                     WPI; 2000-586382/55.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Misc-difference
                                                                                                           proteins they encode, are useful for immunodiagnostic and
                                                                                                                                                                                                           Example 10; Column
                                                                                                                                                                                                                                                                             N-PSDB; A50631
                                                                                                                                                                                                                                                                                                       Stamenkovic I,
                                                                                                                                                                                                                                                                                                                                                                       01-DEC-1992;
25-FEB-1988;
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Misc-difference 305
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Misc-difference 216
                                                                                                                                                                                                                                                                                                                          (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                    28-OCT-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Misc-difference
                                                                                                  mmunotherapeutic applications,
                                                                                                                                                                                                                                    nfections or disorders, e.g. asthma, immune-complex disease,
                                                                                                                                                                                                                                                                                                                                                    89US-0379076
90US-0498809
                                                                                                                                                                                                                                                                                                                                                                       92US-0983647
88US-0160416
                                                                                                                                                                                                                                                                                                       Seed
                                                                                                                                                                                                                                                                                                                                                                                                   98US-0181612
                                                                                                                                                                                                                                                                                                                                             90US-0553759
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338
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306
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268
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220
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                                                                                                                                                                                                         53-55; 75pp;
                                                                                                                                                                                                                                                                                                       В;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          "encoded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "encoded by
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                                                                                                                                                                                                        English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GGT"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAT"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CAA.
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parasitic

surfaces other than macrophages, protein of FcRI and a receptor 1 The present sequence is that of a human macrophage specific PCRI, as deduced from cDNA clone p135 (see A50631), which was isolated from a cDNA library expressed in COS cells using a novel method of the invention designed to isolate CSA nucleic acids. The method is immune-complex disease, amyloidosis, parasitic diseases or multiple sclerosis. FCRI is a high affinity receptor for the FC portion of 19G, normally located on the cell surfaces of macrophages. The ability to interfere with such bonding, or to cause it to occur on the control of the control treatment of immune-mediated infections, diseases, and diso animals, including humans. These disorders include asthma, (panning on) an antibody-coated substrate such as a culture dish. CSA nucleic acids isolated by the method of the invention, and the physical selection of cells expressing the antigen by adhesion to the potencies of antibodies or ligand will in therapy. including the diagnosis and in therapy. be helpful and disorders in The method is to increase

Sequence 344 AA;

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Query Match
Best Local Similarity
----hes 12; Conserv
        Conservative
               100.0%;
       0;
      Score 64; DB ZI;
Pred. No. 0.00025;
                      Length 344;
       Indels
       0;
      Gaps
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1 FYMGSKTLRGRN 12

0;

RESULT
R22550
ID R2
XX
AC R2
XX
DT 21
XX

R22550;

R22550 standard; Protein;

374 AA

21-MAY-1992

(first entry)

Вp Ş

229

240

1 FYMGSKTERGRN 12 fymgsktlrgrn

<u>,</u>

Indels

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RESULT
R20811
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Query Match 100
Best Local Similarity 100
Matches 12; Conservative
                                                                                                                                                                                                                                                                                 typical features of a type I integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane-spanning domain, and a short, highly charged cytoplasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and six Cys residues distributed among three C2 set Ig-related domains. A fusion protein of FCRI and a receptor ligand will be helpful to increase the
                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This amino acid sequence was predicted from the cDNA sequence of cDNA clone p135. It differs from the sequence predicted from clones p98/X2 and p90 (see Q21179 and Q21180, respectively) at position 25; p135 encodes a Ser residue and the other two clone predict a Thr residue. At position 58, p135 predicts Leu and ps predicts Val. Sequences predicted from all 3 clones show the tentile of the control of the con
                                                                                                                                                                                                                                                           potency
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 10; Page 94a; 160pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New CD53 cell surface antigen and DNA encoding immuno-therapy and diagnosis of haematopoietic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Seed B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15-JUL-1990;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO9201049-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rapid immunoselection cloning technique; cell surface antigen; immunodiagnosis; high affinity receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-MAY-1992
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (GEHO-)
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                                                                                                                                                                                                                                                       of antibodies in therapy.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Aruffo A, Amiot M;
                                                                                                                                                                                       374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        90US-0553759
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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                                         100.0%;
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Score 64; DB 13;
Pred. No. 0.00028;
; Mismatches 0;
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                                                                   Length 374;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        clones
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FT FT XXXXXX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 100
Best Local Similarity 100
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cDNA clone p30: It differs from the sequence predicted from clone p135 (see Q21178) at position 25; p135 encodes a Ser residue and clone p90 predicts a Thr residue. At position 58, p90 predicts val while clones p135 and p98/X2 (see Q21179) both predict Leu. Sequences predicted from all 3 clones show the typical features of a type I integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane-spanning domain, and a short, highly charged cytoplasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and six Cys residues distributed among three C2 set Ig-related domains. A fusion protein of FCRI and a receptor ligand will be a close to the contains six potential N-linked glycosylation sites and six Cys residues distributed among three C2 set Ig-related domains. A fusion protein of FCRI and a receptor ligand will be
                                                                                                                                                                                                                           Cell surface antigen; cloning; immunoselection; therapy; diagnosis; vector; FcRI; Fc receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                              R91438
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R91438 standard; Protein; 374 AA
     Misc-difference
                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                             Human FcRI (cDNA clone p135 product).
                                                                                                                                                                                                                                                                                                                                                                 30,-OCT-1996 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This amino acid sequence was
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1992-056864/07
N-PSDB; Q21180.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-JAN-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo, sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rapid immunoselection cloning technique; cell surface antigen; immunodlagnosis; high affinity receptor.
                                                                                    Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         helpful to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 10; Page 94a; 160pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New CD53 cell surface antigen and DNA encoding it - for immuno-therapy and diagnosis of haematopoietic neoplasms,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-JUL-1990;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human macrophage-specific FcRI receptor encoded by clone p90.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (GEHO-) GEN HOSPITAL CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            229
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 FYMGSKTLRGRN 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       fymgsktlrgrn 240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         increase the potency of antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                90US-0553759
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                                                                                 Location/Qualifiers 25
                          /note= "amino acid 25 is Thr in clone p90 and p98
    translated products"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Amiot M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; So 100.0%; P. 100.0%; P. 151ve 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 64; DB 13;
Pred. No. 0.00028;
0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              predicted from the cDNA sequence of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         in therapy.
                                                                                                                                                                                                                                                  immunotherapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 374;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             etc.
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W00859
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                                                                                       XYXYX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 12; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
18-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             for the FC portion of IgG, normally located on cell surfaces of macrophages. Another isolated cDNA clone, p90 (T14719), coded for a variant (W00859) of the FCRI sequence, and a third clone, p98 (T14718), coded for an FCRI (R91439) having a different C-terminal sequence. A fusion protein of FCRI and a receptor ligand will be helpful in increasing the potency of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cloning of cDNA encoding cell surface antigen - useful for isolation of diagnostic and therapeutic proteins % \left( 1\right) =\left\{ 1\right\} 
                                                                                                                                                                                                                     US5506126-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The amino acid sequence (R91438) of human FcRI was detd. from a cDNA clone, p135 (T14717), obtd. from a cDNA library using an immunoselection cloning method. FcRI is a high affinity receptor
                             01-DEC-1992;
25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                 Cell surface antigen; cloning; immunoselection; therapy; diagnosis; vector; FcRI; Fc receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        W00859;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Aruffo A,
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                                                                                                               25-FEB-1988;
                                                                                                                                                                   09-APR-1996
                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                      Human FcRI (cDNA clone p90 product).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            W00859 standard; Protein; 374 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 10; Column 55-56; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1996-200279/20
N-PSDB; T14717.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-DEC-1992
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (GEHO ) GEN HOSPITAL CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
92US-0983647.
88US-0160416.
89US-0379076.
                                                                                                               88US-0160416
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93US-0139273
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89US-0379076
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    translated product"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%;
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Pred. No. 0.00028;
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Length 374; Indels

0,

Gaps

0

immunotherapy;

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RESULT 10
W80448
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Best Local Similarity
Matches 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The amino acid sequence (W00859) of human FCRI was detd. from a CDNA clone, p90 (T14719), obtd. from a cDNA library using an immunoselection cloning method. FCRI is a high affinity receptor for the Fc portion of IgG, normally located on cell surfaces of macrophages. Other CDNA clones (see also T14717-18) coded for variants (R91438-39) of the FCRI sequence. A fusion protein of FCRI and a receptor ligand will be helpful in increasing the potency of antibodies in therapy
 Misc-difference
                     Misc-difference
                                                                Misc-difference
                                                                                                          Misc-difference
                                           Misc-difference
                                                                                     Misc-difference
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                                                                                                                                                                                                                                                                                                                                 07-JUN-1999
                                                                                                                                                                                                                                                                                                                                                      W80448;
                                                                                                                                                                                                                                                                                                                                                                         W80448 standard; Protein; 374 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cloning of cDNA encoding cell surface antigen -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1996-200279/20
N-PSDB; T14719.
                                                                                                                                                                           Misc-difference
                                                                                                                                                                                                                     Misc-difference
                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                   Fc receptor I; FcRI; cell surface antigen; lymphocyte; human;
                                                                                                                                                                                                                                                                                                           Human Fc receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                   · 1 FYMGSKTLRGRN 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 10; Column 55-56; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       of diagnostic and therapeutic proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13-JUL-1990;
18-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                              229 fymgsktlrgrn 240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Seed B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                  /note=
                                                                                                                                                                                               /note=
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93US-0139273
                   /note= "encoded by 141
                                          /note=
103
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99
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77
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                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                           /note=
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0
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Pred. No. 0.00028
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                                                                                              bу
                                                                                                                   bγ
         by GGC"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                CCC.
                                                    CAA"
                                                                          TCC"
                                                                                                AAT"
                                                                                                                     CCC"
                                                                                                                                                                                     CTC"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  useful for isolation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 374;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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expression libraries is based on transient expression of an antigen by adhesion to an antibedy-coated substrate. The method is useful for the isolation and cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V63442-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated genes can be expressed in a prokaryotic or eukaryotic host cells to produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V63444) which allow
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
21-MAY-1997;
Sequence
                                                               the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and immunotherapeutic applications, including the diagnosis and immunotherapeutic applications, including the diagnosis and immunotherapeutic applications.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially human lymphocyte antigenic sequences
                                                    treatment of immune-mediated infections,
                                                                                                                                                                                                                                                                                                     were also obtained.
                                                                                                                                                                                                                                                                                                                                                           cells.
                                                                                                                                                                                                                                                                                                                                                                                         This is the amino acid sequence of human Fc receptor I (FCRI), deduced from cDNA clone pl35 (see V63456) isolated using a rap
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US5830731-A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Misc-difference
                                                                                                                                                                                                                                                                                                 from a single patient undergoing extracorporeal interleukin-2 induction therapy. Clones p90 (see X07372) and p98 (see X07373) were also obtained. A novel method for cloning cDNAs from mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 10; Column 53-54; 75pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-DEC-1992;
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                                                                                                                                                                                                                                                                                                                                                                        immunoselection cloning method from a cDNA library expressed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1998-609251/51
                                                                                                                                                                                                                                                                                                                                                      The cDNA library was constructed from polyA RNA of cells
                                   including humans.
ΑĀ,
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88US-0160416
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176
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RESULT
W97833
ID W97
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AC W97
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Best Local Similarity
Matches 12; Conserva
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                 W97833 standard; Protein; 374 AA
                                                      1 FYMGSKTLRGRN 12
                                                                                       Conservative
                                                                                              100.0%;
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Pred. No.
                                                                                       Mismatches
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                                                                                              .00028;
                                                                                                    Length 374;
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W97833;

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01-DEC-1992;
25-FEB-1988;
                                                      This is the amino acid sequence of human Fc receptor I (FCRI), as deduced from cDNA clone p90 (see X07372) isolated using a rapid immunoselection cloning method from a cDNA library expressed in Cells. The cDNA library was constructed from polyA RNA of cells
                                                                                                                                                      New cloning vector and polylinker - based on existing efficient cloning and expression of mammalian cDNA(s), human lymphocyte antigenic sequences
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Misc-difference 159
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Misc-difference 99
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N-PSDB; X07372.
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              from a single patient undergoing extracorporeal induction therapy. Clones p135 (see V63456) and were also obtained. A novel method for cloning
                                                                                                                            Example 10; Column 53-54; 75pp; English.
                                                                                                                                                                                                                                                         Aruffo A,
                                                                                                                                                                                                                                                                                                                21-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                 21-мау-1997;
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  expression libraries
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85
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and immunotherapeutic applications, including the diagnosis and treatment of immune-mediated infections diseases, and disorders of
                                                                   Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human Fc
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29 fymgsktlrgrn 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     an antibody-coated substrate.
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Matches 12; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-JUL-1989;
13-JUL-1990;
21-MAY-1997;
                          Macrophage; FCRI; cell surface antigen; human; immunoselection; panning; immunodiagnosis; diagnosis; immunotherapy; gene therapy; immune disorder; infection; asthma; immune-complex disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This is the amino acid sequence of human Fc receptor I (FCRI), as deduced from cDNA clone p98/X2 (see X07373) isolated using a rapid immunoselection cloning method from a cDNA library expressed in Cocells. The cDNA library was constructed from polyA RNA of cells
                                                                                       Human macrophage-specific FcRI
                                                                                                                     19-DEC-2000
                                                                                                                                                    Y96134;
                                                                                                                                                                                Y96134 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New cloning vector and polylinker - based on existing sequences fefficient cloning and expression of mammalian cDNA(s), especially
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-DEC-1992;
25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   were also obtained. A novel method for cloning cDNAs from mammalian expression libraries is based on transient expression of an antigen in eukaryotic cells and selection of cells expressing the antigen by adhesion to an antibody coated substrate. The method is useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           from a single patient undergoing extracorporeal interleukin-2 induction therapy. Clones p135 (see V63456) and p90 (see X07372)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Aruffo A,
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DB; X07373.
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89US-0379076.
90US-0553759.
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                                                                                                                                                                                Protein;
              sclerosis;
                                                                                                                                                                                374
                                                                                                                                                                                                                                                                                                                    0,
                                                                                                                                                                                                                                                                                                                 Score 64; DB 19;
Pred. No. 0.00028;
); Mismatches 0;
                                                                                                                                                                                8
            parasitic disease;
                                                                                                                                                                                                                                                                                                                                             Length 374;
                                                                                                                                                                                                                                                                                                                  Indels
            Fc receptor
                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                 01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
23-MAR-1990;
13-JUL-1990;
                         Stamenkovic
                                                                                                                                                                                                                                                                                                                                          Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo
                                                     (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                            28-OCT-1998;
                                                                                                                                                                                                                                       US6111093-A
                                                                                                                                                                                                                                                                                   Misc-difference
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122

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92US-0983647. 88US-0160416. 89US-0379076. 90US-0498809. 90US-0553759.

Seed

98US-0181612

2000-586382/55

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B
                                                                                                                            PENDENDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence is that of a human macrophage specific FCRI, as deduced from cDNA clone p135 (see A50592), which was isolated from a cDNA library expressed in COS cells using a novel method of the invention designed to isolate CSA nucleic acids. The method is based upon transient expression of a CSA in eukaryotic cells and
                                                                                                                                                                                                                                                                                         Human; high affinity Fc receptor; FcgammaRI; immunoglobulin; infection; immune response; CD64; monocyte; macrophage; neutrophil; eosinophil, HIV, IGG; immunosuppressive; antirheumatic; cytostatic; antiinflammatory; AIDS; SLE; multiple myeloma; rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 physical selection of cells expressing the antigen by adhesion to (panning on) an antibody-coated substrate such as a culture dish. CSA nucleic acids isolated by the method of the invention, and the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Isolated nucleic acid molecule encoding the CD19 cell surface antigen, useful for immunodiagnosis and immunotherapy of immune-mediated infections or disorders, e.g. asthma, immune-complex disease, parasitic
                                                                                                                                                                                                                                                                                                                                                                                                                            11-SEP-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Y96226;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  surfaces other than macrophages,
protein of FcRI and a receptor 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         proteins they encode, are useful for immunodiagnostic and immunotherapeutic applications, including the diagnosis and treatment of immune-mediated infections, diseases, and disorders in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 10; Column 53-55; 75pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            diseases
N-PSDB; A27466
                 WPI; 2000-367968/32
                                                                                          03-DEC-1998;
                                                                                                                            03-DEC-1998;
                                                                                                                                                                 07-JUN-2000
                                                                                                                                                                                                   EP1006183-A1
                                                                                                                                                                                                                                                                          systemic lupus
                                                                                                                                                                                                                                                                                                                                                                                       Human high affinity Fc receptor, FcgammaRI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Y96226 standard; Protein; 374 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 immune-complex disease, amyloidosis, parasitic diseases or multiple sclerosis. FCRI is a high affinity receptor for the FC portion of Igo, normally located on the cell surfaces of macrophages. The ability to interfere with such bonding, or to cause it to occur on
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      N-PSDB; A50592
                                                   (PLAC ) MAX PLANCK GES
                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the potencies of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          nimals, including humans.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             229 fymgsktlrgrn 240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 FYMGSKTLRGRN 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                            erythematosus;
                                                                                        98EP-0122969
                                                                                                                            98EP-0122969
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%;
                                                   FOERDERUNG WISSENSCHAFTEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           These disorders include asthma,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 64;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  is useful in therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.00028;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 374;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  to increase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   fusion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
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role in defending the body against infections. First, pathogens are opsonised by serum immunoglobulins. The resulting complex then binds to cells expressing Fc receptors. FcgammaRI molecules are expressed by monocytes and macrophages, but expression can also be induced on neutrophils and eosinophils. Upon Fc receptor activation, immune effector pathways are activated, leading to immune response. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    glycosylation, cancer
                                                                                                                                   may be modified to produce recombinant versions. The recombinant Fc receptor consist only of the extracellular portion of the receptor and are not glycosylated i.e. they do not have transmembrane domains or signal peptides. The recombinant proteins may be used in immunoassays t determine the immune status of patients with chronic diseases of the immune system, e.g. AIDS, systemic lupus erythematosus (SLE), multiple myeloma (MM), or rheumatoid arthritis. In addition, pharmaceutical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel Fc receptor lacking transmembrane domains, a signal peptide, and {\it glycosylation}, useful for diagnosing and treating immune disorders and
                                                                              compositions containing recombinant proteins may be used to treat or prevent autoimmune diseases, allergies or tumours, especially AIDS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FOGARMMARI IS also known as CD64. Fc receptors play an important role in defending the body against infections. First, pathogens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence is the human high affinity Fc receptor, FcgammaRI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Page 26-28; 60pp; English
                                                                                                                                                                                                                                                                                                                                                                    immune effector
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밁 Q RESULT Matches Query Match Best Local (15 , 1 FYMGSKTLRGRN 12 12; Similarity Conservative 100.0%; 0 Score 64; Pred. No. 0.00028; DB 21; Length 374; Indels 0 Gaps 0

Sequence

374

Human cancer associated protein sequence SEQ ID NO:1128. 08-FEB-2001 B43683 standard; Protein; 399 AA (first entry)

vasotropic; antipsoriatic; antianglogenic; gene therapy; inflammation; immune disorder; haematopoletic cell disorder; autolimmune disorder; allergic reaction; graft versus host disease; organ rejection; haemostatic; thrombolytic; cardiovascular disorder; infection; antidiabetic; antiasthmatic; antirheumatic; antiarthritic; antiviral; antiinflammatory; antithyroid; antiallergic; antibacterial; cardiant; diagnosis; Human; neurological disease; dermatological; neuroprotective; thrombolytic; coagulant; nootropic; cancer associated gene; cancer antigen; detection; cytostatic; drug screening proliferative; vulnerary; immunomodulator; cancer;

Homo sapiens

XX S

WO200055350-A1

ΧPN

08-MAR-2000; 2000WO-US05882.

12-MAR-1999; 99US-0124270.

(HUMA-) HUMAN GENOME SCI INC

Rosen CA, Ruben SM;

WPI; 2000-587533/55

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Claim 11; Page 1739-1740; 2352pp; English.
                                                                                                                                 Novel isolated nucleic acids comprising sequences encoding peptides useful for treating or diagnosing e.g. cancer \dot{}
                                                                                                                                                    N-PSDB; C77892
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antithyroid; antialization, antibacterial; antiviral; dermation coal, coapulant; nootropic; antipsoriatic and antiangiogenic. The polynucleotides and coapulant; properties and entry preventing, treating or ameliorating medical conditions and diagnosing pathological conditions. Polynucleotides, conditions and the present conditions and conders by activating or immune collection may be used to treat immune disorders by activating or immune collection, the proliferation, differentiation or mobilisation of immune collection, the presult of the pressent condulate inflammation, condulate haemostatic or thrombolytic activity, modulate inflammation, concers, cardiovascular disorders, neurological disease and bacterial or cural infections. The peptides, nucleotides, antibodies, agonists and conductions and bacterial or concers, cardiovascular disorders, neurological disease, agonists and salso be used in drug screens. C78449 to C78457 and concerns and conductions are peptides and conductions are peptides. C77607 to C78448 encode the human cancer associated proteins given in B43398 to B44239. The proteins can have activities based on the tissues and cells the genes are expressed in. Example of activities include: cytostatic; proliferative; vulnerary; immunomodulator; antidiabetic; cytostatic; proliferative; vulnerary; immunomodulator; antidiabetic; antiasthmatic; antirheumatic; antiarthritic; antiinflammatory;

Sequence 399 AA;

QΥ Query Match 100.0%; Score 64; DB 21; Best Local Similarity 100.0%; Pred. No. 0.0003; Matches 12; Conservative 0; Mismatches 0; 1 FYMGSKTLRGRN 12 Length 399; Indels 0; Gaps 0;

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Search completed: June 4, 2001, 12:13:29 Job time: 202 sec

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OM protein - protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Database
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Searched
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score
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length: 2000000000
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Gapop 10.0 , Gapext 0.5
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64
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                                          1 FYMGSKTLRGRN, 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /cgn2_6/ptodata/2/iaa/5A_COMB.pep:*
/cgn2_6/ptodata/2/iaa/5B_COMB.pep:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /cgn2_6/ptodata/2/iaa/6A_COMB.pep:*
/cgn2_6/ptodata/2/iaa/6B_COMB.pep:*
/cgn2_6/ptodata/2/iaa/PCTUS_COMB.pep:*
/cgn2_6/ptodata/2/iaa/backfiles1.pep:*
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                                                                                        365
365
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4 US-08-061-376-5
1 US-08-403-866-5
1 US-08-403-866-5
1 US-08-403-866-5
1 US-08-428-414A-2
2 US-08-428-418-438-4
2 US-08-427-4978-9
4 US-09-405-035-2
1 US-09-405-035-3
1 US-09-405-035-4
9CT-US94-09789-2
US-08-560-098A-45
US-08-560-098A-45
US-08-560-098A-45
US-08-560-098A-45
US-08-560-098A-45
US-08-967-024C-24
US-08-967-024C-24
US-08-967-024C-25
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US-08-080-255-7
US-08-465-713-7
PCT-US93-05857-7
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Sequence
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31	<u>3</u> 1	31	31	31	31	31	32	32	32	32	32	32	32	32	32	32	32
48.4	48.4	48.4	48.4	48.4	48.4	48.4	50.0	50.0	50.0	50.0	50.0	50.0	50.0	50.0	50.0	50.0	50.0
474	383	260	260	2601	260	70	570	570	570	570	561	561	561	432	431	430	430
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US-08-961-083-60	US-08-464-523B-31	US-09-012-431-139	US-08-639-075A-139	US-08-906-616-139	us-08-906-769-139	US-09-003-708A-4	US-09-368-408-7	US-09-368-408-1	US-08-967-364-7	US-08-967-364-1	US-08-693-835-2	US-08-566-347-2	US-08-358-901-2	US-08-560-098A-47	5188829-1	5219569-2	US-07-942-157A-3
Sequence 60, Appl	Sequence 31, Appl	Sequence 139, App	Sequence 139, App	Sequence 139, App	Sequence 139, App	Sequence 4, Appli	Sequence 7, Appli	. Sequence 1, Appli	Sequence 7, Appli	Sequence 1, Appli	Sequence 2, Appli	Sequence 2, Appli	Sequence 2, Appli	Sequence 47, Appl	Patent No. 5188829	Patent No. 5219569	Sequence 3, Appli

ALIGNMENTS

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TOPOLOGY:
US-08-332-562A-94
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                                                                                                                          APPLICATION NUMBER: .US 07/896,457
FILING DATE: 27-MAY-1992
ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 54270/119/GRHA
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
TELEPAX: (202)672-5399
                                                                                                  INFORMATION FOR SEQ ID NO:
                             SEQUENCE CHARACTERISTICS:
LENGTH: 91 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                   CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Rel CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                      FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY:
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                                                                                                                904136
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             USA
                                                                                                                                                                                                                                                                                                                                                                      PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HULETT, Mark D.

ENTION: FC RECEPTOR FOR IMMUNOGLOBULIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HOGARTH, Mark
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Margaret L
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Best Local Similarity

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US-08-465-713-7
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US-08-080-255-7
                                                                                                                                Sequence 7, Application US/08465713 Patent No. 6121419
                                                                                                                                                                                                                                                                                               Matches
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                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: DNA (genomic) -08-080-255-7
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                                                                                                                ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (512) 474-7577 NFORMATION FOR SEQ ID NO:
                             TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                            APPLICANT: Rowley, Janet E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Parker, David L.
REGISTRATION NUMBER: 32,165
REGISTRATION NUMBER: AR
REFERENCE/DOCKET NUMBER: AR
TELECOMMUNICATION INFORMATION:
TELEPHONE: (512) 320-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/080,255
FILING DATE: 19930617
                  NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STATE: Town
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ATTORNEY/AGENT INFORMATION:
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ORRESPONDENCE ADDRESS:
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                                                                                               Rowley, Janet D.
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                                                               COMPOSITIONS AND METHODS FOR
                              DETECTING GENE REARRANGEMENTS AND TRANSLOCATIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DETECTING GENE REARRANGEMENTS AND TRANSLOCATIONS
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1; Mismatches
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Best Local S
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INFORMATION FOR SEQ ID NO:
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REFERENCE/DOCKET NUMBER: ARCD
TELECOMMUNICATION INFORMATION:
TELEPHONE: (512) 320-7200
TELEPAX: (512) 474-7577
NEORMATION TO TELEPAX: (512) 474-7577
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 0
FILING DATE: 17/06/92
                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/465,713
FILING DATE: 06-UUN-1995
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MEDIUM TYPE: Floppy disk
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APPLICATION NUMBER: US/08/080,255
FILING DATE: 17 JUNE 1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
                                                                                                    APPLICATION NUMBER: PFILING DATE: 19930617 CLASSIFICATION:
                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0,
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OPERATING SYSTEM:
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THE UNIVERSITY OF TEXAS SYSTEM
TO UNIVERSITY OF TEXAS TO TRANSLOCATIONS
TO THE UNIVERSITY OF TEXAS TO THE UNIVERSITY OF TEXAS TO THE UNIVERSITY OF TEXAS TO THE UNIVERSITY OF TEXAS TO THE UNIVERSITY OF TEXAS TO THE UNIVERSITY OF TEXAS TO THE UNIVERSITY OF TEXAS TO THE UNIVERSITY OF TEXAS TO THE UNIVERSITY OF TEXAS TO THE UNIVERSITY OF TEXAS TO THE UNIVERSITY OF TEXAS TO THE UNIVERSITY OF TEXAS TO THE UNIVERSITY OF TEXAS TO THE UNIVERSITY OF TEXAS TO THE UNIVERSITY OF TEXAS TO THE UNIVERSITY OF TEXAS TO THE UNIVERSITY OF TEXAS TO THE UNIVERSITY OF TEXAS TO THE UNIVERSITY OF TEXAS TO THE UNIVERSITY OF THE UNIVERSITY OF TEXAS TO THE UNIVERSITY OF THE UNIVERSITY OF THE UNIVERSITY OF TEXAS TO THE UNIVERSITY OF THE U
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SYSTEM: PC-DOS/MS-DOS
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PCT-US93-05857-7
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INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                     TELEFAX: (619)546-939
NFORMATION FOR SEQ ID NO:
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LENGTH: 1400 amino acids
TYPE: amino acid
                                                                                                                                                                      SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Patentin Rele
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                 TELEPHONE: (619)546-4737
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: 444 SOUTH
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                           NAME: Reiter, Stephen E.
REGISTRATION NUMBER: 31,192
REFERENCE/DOCKET NUMBER: P4
                                                                                                               TOPOLOGY:
                                                                                                                          STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM:
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REFERENCE/DOCKET NUMBER: ARCD:072/PAR
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REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   E OF INVENTION: CHARACTERIZATION OF A CHROMOSOME 11Q23
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                                                                                                                                            amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             E: Pretty, Schroeder, Brueggemann & Clark 444 South Flower Street, Suite 2000
                                                                                                                                                      3969 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                    USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Djabali, Malek
Selleri, Licia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (512)
                                                                                                                                                                                                  (619)546-9392
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Parry, Pauline
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Evans, Glen A
                                                                                                                                                                                                                                                                                                                                                                                          IBM PC compatible
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66.7%;
                           57.8%;
66.7%;
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                                                                                                                                                                                      5:
                         Score 37;
Pred. No.
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Pred. No.
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                         1.6e+02;
                                          DB 4;
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                                       Length 3969;
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            Indels
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          Gaps
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                                                                                                                                                                                                                                                     RESULT
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                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 70.0
Matches 7; Conservative
                                                                                                                                                                                                    Sequence 7, Application US/08469667 Patent No. 5733748
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEX: 978450 (WUT)
INFORMATION FOR SEQ ID NO: 5:
                                                                                                          NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS
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ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FITLE OF INVENTION:
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                                                                                                                                           TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                  129 MGAKSLIGRN 138
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                                            CITY: Roseland
                                                            STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Lactococcus lactis subsp. lactis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Goldman, Michael L. REGISTRATION NUMBER: 30,
                COUNTRY:
                                                                             ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                               3 MGSKTLRGRN 12
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                                                                                           DDRESSEE:
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07068-1739
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    978450
                                                              6 Becker Farm Road
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                USA
                                                                                                                                                        Rosen, Craig
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Godon, Jean-Jacques
                                                                                                                                                                       Yu, Guo-Liang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ehrlich, Stanislav
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         : Floppy disk
IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         linear
                                                                             Carella, Byrne, Bain, Gilfillan, Cecchi
Stewart & Olstein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (716) 263-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nucleic acid coding for an alpha-acetolactate synthase from Lactococcus and its applications
                                                                                                                                         Colon Specific Genes and Proteins
                                                                                                                                                                                                                                                                                                                                                                               56.2%;
70.0%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                               DB 1;
11;
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COMPUTER READABLE FORM:

<u>ა</u>

Length 135;

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PCT-US95-07289-7
CT-US95-07289-7
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GENERAL INFORMATION:
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                                                                      TELEFAX: 201-994-1/44
NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                 TELECOMMUNICATION INFORMATION: TELEPHONE: 201-994-1700
            MOLECULE TYPE: protein
                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: FETTATO, GTEGOTY D.
REGISTRATION NUMBER: 36,134
REFERENCE/DOCKET NUMBER: 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 135 amino acids
                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Rosen, Craig
TITLE OF INVENTION: Colon Specific Genes and Proteins
NUMBER OF SEQUENCES: 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET:
                               TOPOLOGY:
                                                                                                                                                                                                                                                       APPLICATION NUMBER: FILING DATE: 06-JU
                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                     CLASSIFICATION:
                                                                                                                                                                                                                                                                                                        SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 07068-1739
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Ferraro, Gregory D.
REGISTRATION NUMBER: 36,134
REFERENCE/DOCKET NUMBER: 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/469,667-
FILING DATE: 06-JUN-1995
                                                                 LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
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                                            amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                               Roseland
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                                                             135 amino acids
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E: Stewart & Olstein
6 Becker Farm Road
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                                                                                                                                                                                                                                                                                                       Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Yu, Guo-Liang
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                                                                                                                                                                                                                                                       06-JUN-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    51.6%;
45.5%;
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Best Local Similarity
"hes 6; Conserv
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                                                                                                                                                   RESULT 10
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                                                                                                   Sequence 4, Application US/08341843B Patent No. 5872225
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
5; Conserv
                          Patent No. 5872225
                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patent No. 5912166
                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (206) 682-6031
TELEX: 3723836 SEEDANDBERRY
INFORMATION FOR SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                       TITLE OF INVENTION:
                                                                          APPLICANT:
                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: SEED and BERRY
STREET: 6300 Columbia Cen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: 436
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                             51 FIMGKKTLQAK 61
                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/0 FILING DATE: 21-APR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                           LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                           amino acid
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Kadlecek, Ann T.
Kadlecek, 39,244
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Application US/08428414A
                                                                                                                                                                                                                                                                                                                                                                                           322 amino acids
                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                               linear
                                      Vance
A Method for Characterizing the
Nucleotide Sequence of LICAM and
                                                                                                                                                                                                                                                                      51.6%;
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the Nucleotide Sequence
Characterized Thereby
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Pred. No.
                                                                                                                                                                                                                                                                      Score 33; DE
Pred. No. 58;
                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     701 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB
22;
                                                                                                                                                                                                                                                                                   Length 322;
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UMBER OF SEQUENCES: 3

ADDRESSEE: Fay, Sharpe, Bea ADDRESSEE: Minnich & McKee

Fay, Sharpe, Beall, Fagan,

STREET:

Suite 700

1100 Superior Avenue

CITY: Cleveland

Ohio

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                                                                        Query Match
Best Local
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MEDIUM TYPE: Diskett
MEDIUM TYPE: storabl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NOLECULE TYPE: amino acids YPOTHETICAL: irrelevant
                                                                                                                                                                                                                                                                                                                                                           CLONE: synthesis of several clones UBLICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ELECOMMUNICATION INFORMATION: TELEPHONE: (216) 861-5582
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 07/9
FILING DATE: June 26, 19
TTORNEY/AGENT INFORMATION:
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56 YSGSFTIEGNN 66
                                                      Local Similarity es 6; Conserv
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SOFTWARE: ASCII
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS:
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                            2 YMGSKTLRGRN 12
                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: mouse
INDIVIDUAL ISOLATE: 8 day old mouse brain
                                                                                                                                                                                                                                                                                                                                                                                                                                                              I-SENSE:
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44114-2518
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1988
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(216) 98
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                                                        Conservative
                                                                                                                                                                                                                                                                                                  Teplow, D.
                                                                                                                                                                                                                                                                                                                                            MOOS, M.
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                                                                    Score 32;
Pred. No.
                                                        Mismatches
                                                                     DB 2;
23;
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JS-08-427-497E-9
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FILING DATE: April 24, 1995
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MEDIUM TYPE: storable
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701-703
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                                                                                        Neural adhesion molecule L1 is a member of the immunoglobulin
                                                                      superfamily
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1100 Superior Avenue
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A Method for Characterizing the
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; DATE: 1988 US-08-427-497E-9

Query Match

50.0%; 54.5%;

Score 32; Pred. No.

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Best Local Similarity 66.7
Matches 6; Conservative
                                           Matches
                                                            Query Match
Best Local
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Matches 6; Conservative
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APPLICANT: Biere, Anja Leona
APPLICANT: Citron, Martin
                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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Patent No. 6184351
                                                                                                                                     ORGANISM: ADULT HUMAN BRAIN -09-405-035-2
                                                                                                                                                                                                                            SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/405,035
CURRENT FILING DATE: 1999-09-24
EARLIER APPLICATION NUMBER: 60/101,862
EARLIER FILING DATE: 1998-09-25
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NUMBER OF SEQ ID NOS: 6
SOFTWARE: Patentin Ver. 2.0
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CURRENT FILING DATE: 1999-09-24
                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: ALPHA-SYNUCLEIN SUPER-MUTANTS ACCELERATE ITLE OF INVENTION: ALPHA-SYNUCLEIN AGGREGATION THE REFERENCE: A-565
                                                                                                                                                                                                                                                                                                                                                                                        PPLICANT: Biere, Anja Leona
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TYPE: PRT
                                                                                                                                                                         TYPE: PRT
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                                                                                                                                                                                        ENGTH: 140
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       2 YMGSKTLRG 10
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66.7%;
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Pred. No. 36;
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Search completed: June Job time: 223 sec

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; LENGTH: 140
; TYPE: PRT
; ORGANISM: ADULT HUMAN BRAIN US-09-405-035-4
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; TYPE: PRT
; ORGANISM: ADULT HUMAN BRAIN
US-09-405-035-3
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Patent No. 6184351
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CURRENT FILING DATE: 1999-09-24
EARLIER APPLICATION NUMBER: 60/101,862
EARLIER FILING DATE: 1998-09-25
NUMBER OF SEQ ID NOS: 6
                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: ALPHA-SYNUCLEIN SUPER-MUTANTS ACCELERATE TITLE OF INVENTION: ALPHA-SYNUCLEIN AGGREGATION FILE REFERENCE: A-565
CURRENT APPLICATION NUMBER: US/09/405,035
CURRENT FILING DATE: 1999-09-24
CURRENT FILING DATE: 1999-09-24
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EARLIER FILING DATE: 1998-09-25
NUMBER OF SEQ ID NOS: 6
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APPLICANT: Citron, Martin
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APPLICANT: Citron, Martin
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39 YVGSKTKEG
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Pred. No. 36;
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score
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Match Length DB
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Fc gamma (IgG) rec
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408	383	379	377	335	291	278	267	249	236	167	165	107	771	736	531
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T18813	A41545	A82806	A44216	G64385	S10867	TPHUTW	D72499	S75500	T16164	T29565	E70486	T16521	H72410	S57961	S76041
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ALIGNMENTS

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Science
                                      A;Reference number: A64720; MUID:97426617
A;Accession: H64726
                                                                                                                                       R;Blattner, F.R.; Plunkett III, G.; Bloch, C.A.; Perna, N.T.; Burland, V.; Riley,
                 A;Status: nucleic acid sequence not shown;
                                                                                                                                                                                                                              yabP protein
                                                                                                                                                                 Species: Escherichia coli
;Date: 12-Sep-1997 #sequence_revision 17-Sep-1997 #text_change 04-Mar-2000
;Accession: H64726
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ;Map position:
                                                                             ;Title: The complete genome sequence of Escherichia
                                                                                                                                                               Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ;Molecule type: mRNA;
;Residues: 1-24,'S',26-337,'T',339-374 <AL2>;
;Residues: 1-24,'S',26-337,'T',339-374 <AL2>;
;Cross-references: EMBL:X14356; NID:g31331; PIDN:CAA32537.1; PI;Perez, C.; Wietzerbin, J.; Benech, P.D.
ol. Cell. Biol. 13, 2182-2192, 1993
ol. Cell. Biol. 13, 2182-2192, 1993
;Title: Two cis.DNA elements involved in myeloid-cell-specific
Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ucleic Acids Res. 16,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Superfamily: Fc gamma receptor I; immunoglobulin homology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cross-references: GB:S57204; NID:g298692;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Reference number: Accession: .I57525
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Title: Nucleotide sequence of three cDNAs Reference number: S03018; MUID:89098339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cross-references: GB:L03418; Allen, J.M.; Seed, B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Residues: 1-374 <RES>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Accession: I70304
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Keywords: alternative splicing; glycoprotein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cross-references: GDB:135911; OMIM:146760
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gene: GDB:FCGR1A; CD64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Molecule type: DNA
Residues: 1-8 <RE2>
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Accession: B41357
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293-313/Domain: transmembrane #status predicte
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                                                                                                                          ; Rose,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  l6-292/Domain: extracellular #status predicted <EXT>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                translated from
                                                                                                   D.J.; Mau, B.; Shao, Y.
7, 1453-1462, 1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      translated from GB/EMBL/DDBJ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Redecha, P.B.;
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                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                            Score 64; DB 1; Le
Pred. No. 7.5e-05;
Pred. matches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      carbohydrate
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                   translation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        <SIG>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  in human mononuclear cells.
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C; Superfamily: Fc gamma
C; Keywords: immunoglobul
                                                                                                                                                                                                                                                                                                                                                                                                                N;Alternate names: high affinity IgG receptor
C;Species: Mus musculus (house mouse)
C;Date: 18-Jun-1993 #sequence_revision 18-Nov-1994 #text_change
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Fc gamma (IgG) receptor high affinity C;Species: Mus musculus (house mouse) C;Date: 02-Jul-1996 #sequence_revision
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A:Experimental sourc
C:Genetics:
                A;Cross-references: GB:M31314; NID:g200752; PIDN:AAA40056.1; C;Superfamily: Fc gamma receptor I; immunoglobulin homology
                                                          A;Residues:
                                                                                             A; Status: preliminary
                                                                                                                                     A; Title: Molecular cloning and expression of A; Reference number: A43511; MUID:90111035
                                                                                                                                                                                                                                                                                                          3;Osman, N.; Kozak, ......
J. Immunol. 148; 1570-1575, 1992
A;Title: Structure and mapping of the gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Fc gamma (IgG) receptor high affinity - mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       F;128-180/Domain:
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C;Keywords: immunoglobulin receptor
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A; Residues: 1-336 < RES>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 R; Prins, J.B.;
Science 260, 6
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                                                                             A; Molecule type: mRNA
                                                                                                                    A; Accession: A43511
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                                                                                                                                                                                                                                                                                 \;Status: preliminary
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                                                                                                                                                                                              ;Sears, D.W.; Osman, N.;
                                                                                                                                                                                                                                       ;Residues:
                                                                                                                                                                                                Note: sequence extracted from NCBI backbone Sears. D.W.; Osman, N.; Tate, B.; McKenzie,
                                                                                                                                                                                                                                                            ; Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                                                    ;Accession: A46480;
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Best Local
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Best Local
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                                                        1-404 <SEA>
                                                                                                                                                                                                                                       1-404 <OSM>
                                                                                                                                                                                                                                                                                                                                                                               Kozak, C.A.; McKenzie, I.F.; Hogarth,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
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95-698, 1993
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ce: strain K-12, substrain MG1655
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        62.5%;
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54.5%;
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Pred. No. 2.4;
3; Mismatches
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Pred. No. 3.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           02-Jul-1996 #text_change 21-Jan-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       mouse
                                                                                                                                                                                                                                                                                                                                       encoding mouse high affinity Fc
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2.4;
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                                                                                                                                                                                                (NCBIN:85205, I.F.C.; Hogart
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                                                                                                                                                                                                Hogarth,
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                                                                                                                                                                                                                  NCBIN:85208,
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                                 PID:g200753
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C;Date: 14-Aug-1998 #sequence_revision 14-Aug-1998 #text_change 05-Nov-1999
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A;Accession: H71147
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A; Residues: 1-379 < KAW>
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A;Accession: E71162
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DNA Res. 5, 55-76, 1998
                                                                                                                                                                                             A; Experimental source:
                                                                                                                                                                                                      A;Cross-references: GB:AP000002; NID:g3236129; PIDN:BAA29469.1;
                                                                                                                                                                                                                         A; Residues:
                                                                                                                                                                                                                                        A; Molecule type: DNA
                                                                                                                                                                                                                                                       A; Status: preliminary; nucleic
                                                                                                                                                                                                                                                                                                                                    R;Kawarabayasi, Y.; Sawada, M.; Horikawa, H.; Haikawa, M.; Ohfuku, Y.; Funahashi, T.; Tanaka, T.; Kudoh, Y.;
                                                                                                                                                                                                                                                                                                                                                                       C; Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                      hypothetical protein PH0394 - Pyrococcus horikoshii
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                                                                                                                                                                                                                                                                                                                       M.; Ohfuku, Y.; Funaha
NA Res. 5, 55-76, 1998
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;Accession: E71162
                                                               Matches
                                                                            Query Match
Best Local
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Best Local
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nes 7; Conser
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7; Conserv
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8; Conser
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273
                                                                                                                                                                           replaces an interim accession
                                                                                                                                                                                            strain OT3
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Pred.
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Pred. No. 6
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4.7;
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                                                                                                                                                                         for a sequence replaced by GenBank
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            for a sequence replaced by
                                                                                                                                                                                                                                                                                                                                    Y.; Hino, Y.;
Yamazaki, J.;
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Kushida,
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C;Date: 30-Apr-1993 #sequence_revision 18-Nov-1994 #text_change 01-Dec-2000 C;Accession: A44265; A44264; I58112; I37165; I38485 R;Tkachuk, D.C.; Kohler, S.; Cleary, M.L.
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Best Local Similarity
Thehes 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R:Chung, S.O.; Kubo, T.; Natori, S.
J. Biol. Chem. 270, 4624-4631, 1995
A;Title: Molecular cloning and sequencing of arylphorin-binding protein in A;Reference number: A56097; MUID:95181458
A;Accession: A56097
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Title: The complete genome sequence of Escherichia coli K-12 A;Reference number: A64720; MUID:97426617 A;Accession: F64940
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F64940
                                                                                                    A; Reference number: A44265;
A; Accession: A44265
                                                                                                                              A; Title: Involvement of a homolog of Drosophila trithorax A; Reference number: A44265; MUID:93046667
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Molecule type: mRNA
A; Residues: 1-1163 < CHU>
A; Cross-references: GB: D29741
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C;Species: Sarcophaga peregrina
C;Date: 03-Oct-1995 #sequence_revision
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C;Species: Sarcophaga peregrina
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A;Experimental source: strain K-12, substrain MG1655
C;Superfamily: hypothetical protein b1798
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Molecule type: DNA
A; Residues: 1-212 <BLAT>
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                                            A; Molecule type:
A; Residues: 1-39
                                                                                    A; Status: nucleic acid sequence not shown; not compared with conceptual translation
                                                                                                                                                                             Cell 71, 691-700, 1992
                                                                                                                                                                                                                                                                      C:Species: Homo sapiens (man)
                                                                                                                                                                                                                                                                                        N;Alternate names: acute lymphoblastic leukemia gene 1 protein (ALL-1)
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                                                                                                                                                                                                                                                                                                                trithorax homolog HTX, version 2 - human
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Cross-references: GB:L04284; NID:g184393; PIDN:AAA58669.1; PID:g18439; Note: sequence extracted from NCBI backbone (NCBIP:117729)
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8; Conserv
                                            1-3968 <TKA>
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53.3%;
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Y.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         . 55;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 1163;
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A; Molecule type: mRNA
A; Residues: 1317-1700, 'DD', 1703-1936, 'H', 1938-2180, 'S', 2182-2328 <DJA>
A; Residues: 6B:L01986; NID:9307522; PIDN:AAA92511.1; PID:9553800
A; Cross-references: GB:L01986; NID:9307522; PIDN:AAA92511.1; PID:9553800
A; Marschalek, R.; Greil, J.; Lochner, K.; Nilson, I.; Siegler, G.; Zweckt
T. J. Haematol. 90, 308-330, 1995
A; Title: Molecular analysis of the chromosomal breakpoint and fusion tran
Accession: 137165; MUID:95315013
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Cross-references: EMBL:X83604; NID:g897757; PIDN:CAA58584.1; PID:g899268
A;Note: submitted to the EMBL/GenBank/DDBJ databases by R. Marschalek, 20 I
R;Cu, Y.; Alder, H.; Nakamura, T.; Schichman, S.A.; Prasad, R.; Canaani, O
Cancer Res. 54, 2327-2330, 1994
                                                                                                  leuD 3'-region hypothetical protein - Lactococcus lactis subsp. lactis (str
C;Species: Lactococcus lactis subsp. lactis
C;Date: 03-Oct-1995 #sequence_revision 03-Oct-1995 #text_change 17-Mar-2000
                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Thehes 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Status: not compared with conceptual translation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11 71, 701-708, 1992
Title: The t(4;11) chromosome translocation of human Reference number: A44264; MUID:93046668
Accession: A44264
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cross references: GB:L04731; NID:9339921
Note: sequence extracted from NCBI backbone (NCBIP:117779)
R;Godon, J.J.; Delorme, C.; Bardowski, J.; Chopin, M.C.; Ehrlich, S.D.; J. Bacteriol. 175, 4383-4390, 1993
A;Title: Gene inactivation in Lactococcus lactis: branched-chain amino a A;Reference number: A36889; MUID:93322316
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   , Map position:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ;Molecule type: DNA
;Residues: 1251-1486,'G',1488-1538 <RES>
;Cross-references: EMBL:U04737; NID:9451
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ture Genet.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Title: Sequence analysis of the breakpoint cluster region in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Reference number:
Accession: I58112
                                                                                                                                                                                                                                                                                                                                                                                                                                       1479-1506/Region: zinc finger
1527-1556/Region: zinc finger
1528-1596/Region: zinc finger
16873-1990/Region: zinc finger
1933-1995/Region: zinc finger
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Residues: 1212-1603, 'GTE' <MAR>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cross-references:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gene: GDB:MLL; HTX;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        keywords: acute lymphoblastic leukemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Note: the list of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    itle: A trithorax-like gene is interrupted by chromosome 11q23 eference number: I58112; MUID:93265134
                                                                                Accession: F36889
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            idues: 63-316, 'GLLINSELEK', 327, 'Q', 329, 'VR', 332, 'DKEGTPP', 340, LQIESTSP', 2555-3554, 'N', 3556-3594, 'V', 3596-3899, 'A', 3901-3968
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1992
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Pred. No. 2e+0
1; Mismatches
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2e+02;
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                                     amino acid biosynthesis
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, H.;
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                     δÃ
                                                               Query Match
Best Local Similarity
Matches 6; Conserv
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C;Superfamily: unassigned ATP-binding cassette proteins;
C;Keywords: ATP; P-loop
F;19-219/Domain: ATP-binding cassette homology <ABC>
F;36-43/Region: nucleotide-binding motif A (P-loop)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Molecule type: DNA
A; Residues: 1-259 <GOD>
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A; Status: preliminary;
A; Map position: 3
A; Introns: 19/1;
A; Note: C0410w
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              hypothetical protein (leuD 3' region) - Le
C;Species: Lactococcus lactis subsp. lacti
C;Date: 31-Dec-1993 #sequence_revision 31-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Cross-references: EMBL:M90761; NID:g2565137; PIDN:AAB81917.1; PID:g2565155
C;Superfamily: unassigned ATP-binding cassette proteins; ATP-binding cassette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         J. Bacteriol. 174, 6580-6589, 1992:
A;Title: Branched-chain amino acid biosynthesis
A;Reference number: S35132; MUID:93015710
A;Accession: S35136
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 F;19-219/Domain: ATP-binding cassette homology <ABC>F;36-43/Region: nucleotide-binding motif A (P-loop)
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A; Residues: 1-259 <GOD>
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Best Local S
Matches 7
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                                                                                                                                                                          R;Lawson, D.; Bowman, S.; I submitted to the EMBL Data
                                                                                                                                                                                                                             hypothetical protein C0410w - malaria parasite (Plasmodium falciparum) C;Species: Plasmodium falciparum C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 09-Jun
                                                                                                                                                                                                                  C; Accession: T18443
                                                                                                                        A; Status: preliminary; translated from GB/EMBL/DDBJ
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Best Local Similarity 70.0
                                                                       A;Cross-references:
                                                                                             A;Molecule type: I
A;Residues: 1-280
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                                                                                          A;Residues:
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                                                        Genetics:
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7; Conserv
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                    61/3; 87/2; 117/2; 145/3; 171/2
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                                                                           EMBL: Z98547; NID: e1325376; PID: e1407850;
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70.0%;
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70.0%;
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Pred. No.
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Pred. No.
                                                                                                                                                                        , В.
Y, August 1997
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18;
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FYMGSKTLRGRN 12

56.2%;

Score 36; Pred. No.

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Length 280;

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A;Experimental source: cultivar Williams
C;Superimmily: GTP-binding regulatory protein Gs alpha chain
C;Keywords: GTP binding; signal transduction
                                                                                                                                                                                                                                                                                                                                                                                                             GTP-binding regulatory protein alpha chain 2 - soybean
C.Species: Glycine max (soybean)
C.Date: 30-Apr.1999 #sequence_revision 30-Apr.1999 #text_change 11-May-2000
                                                                                                                                                                                                                                                                                                   C; Accession: T07159

R;Kim, W.Y.; Cheong, N.E.; Lee, D.C.; Je, D.Y.; Bahk, J.D.; Cho, M.J.; Lee, S.Y.

Plant Physiol. 108, 1315-1316, 1995

A;Title: Cloning and sequencing analysis of a full-length cDNA encoding a G protein alph

A;Reference number: Z15966; MUID:95357428
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C;Species: Escherichia coli
C;Date: 05-Mar_1995 #sequence_revision 12-May-1995 #text_change 11-Jan-2000
                                                                                                                                                                                                                                                                                 N;Status: preliminary; translated from GB/EMBL/DDBJ
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                                                                  Query Match
Best Local Similarity 58.:
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Residues: 1-312 <TAV>
Cross references: EMBL:x82430; NID:g572684; PID:g572685
                                                                                                                                                                                                                                            Molecule type: mRNA
Residues: 1-385 <KIM>
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2; Mismatches
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P39220; P75632;
D1-FEB-1995 (Re)
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                                 SERQUENCE FROM N.A.
STRAIN-K12 / MG1655;
MEDLINE-97426617; PubMed-9278503;
Blattner F.R., Plunkett G. III, Bloch C.A.,
Riley M., Collado-Vides J., Glasner J.D., Ro
Gregor J., Davis N.W., Kirkpatrick H.A., Goe
Gregor J., Shao Y.;
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              "The complete genome sequence of Escherichia coli K-12.", Science 277:1453-1474(1997).
                                                                                                                      Yura T., Mori H., Nagai H., Nagata T., Ishihama A., Fujita N., Isono K., Mizobuchi K., Nakata A.; "Systematic sequencing of the Escherichia coli genome: analysis the 0-2.4 min region.";
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lin domain; Alternative splicing; Polymorphism.
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                                                                                                        region.";
Res. 20:3305-3308(1992).
                                                                                                                                                                                                                                                                                               STANDARD;
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r J.D., Rode C.K., Mayhew G.F.,
H.A., Goeden M.A., Rose D.J.,
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                                                                                                    the European Bioinformatics Institute. The use by non-profit institutions as long modified and this statement is not removed. entitles requires a license agreement (See or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OSMAN N., KOZAK C.A., MCKENZIE I.F., HOGATH P.M.;
"Structure and mapping of the gene encoding mouse high affinity
gamma RI and chromosomal location of the human Fc gamma RI gene
J. Immunol. 148:1570-1575(1992).
-!- FUNCTION: BINDS TO THE FC REGION OF IMMUNOGLOBULINS GAMMA. I
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EMBL; AE000116; AAC73167.1; -
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Hypothetical protein.
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                                                                                                                                                                                                                                                                                                                                                                                                        -i- SUBCELLULAR LOCATION: TYPE I MEMBRANE PROTEIN.
-i- TISSUE SPECIFICITY: MACROPHAGE-SPECICIC.
-i- SIMILARITY: CONTAINS, 3 IMMUNOGLOBULIN-LIKE C2-TYPE DOMAINS.
-i- CAUTION: IT IS UNCERTAIN WHETHER MET-1 OR MET-10 IS THE INITIATOR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HIGH AFFINITY IN RI) (FCRI) (IGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=92166399; PubMed=1531670;
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IY IMMUNOGLOBULIN GAMMA FC RECEF
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                                                                                                                                               (See http://www.isb-sib.ch/announce/
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InterPro;
                                                                              modified and this statement is not removed. entities requires a license agreement (See
                                                                                                                                               "Isolation, characterization and analysis of the expression of the Leishmania ribosomal po protein genes.";
MOI. Blochem. Parasitol. 61:265-274(193).
-i- FUNCTION: RIBOSOMAL PROTEIN PO IS THE FUNCTIONAL EQUIVALENT OF E.COLI PROTEIN L10.
-i- SUBUNIT: PO FORMS A PENTAMERIC COMPLEX BY INTERACTION WITH DIMERS OF PI AND P2.
-i- SIMILARITY: BELONGS TO THE L10P FAMILY OF RIBOSOMAL PROTEINS.
                                      EMBL; X72714; CAA51264.1; -. EMBL; X72714; CAA51263.1; -.
                                                                    or send an
                                                                                                                   This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss_Institute_of Bioinformatics and the EMBL outstation -
                                                                                                                                                                                                                                                      STRAIN-LEM 75
                                                                                                                                                                                                                                                                                 Eukaryota; Euglenozoa;
NCBI_TaxID=5671;
                                                                                                                                                                                                                                                                                                                 60S ACIDIC RIBOSOMAL PROTEIN PO. LIPO-A AND LIPO-B.
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P39097;
01-FEB-1995 (Rel.
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                                                                                                           the European Bioinformatics Institute.
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Ribosomal_L10;
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IG-LIKE C2-TYPE DOMAIN.
IG-LIKE C2-TYPE DOMAIN.
N-LINKED (GLCNAC.))(
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Pred. No.
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P76249; 007971; 007969;
15-JUL-1998 (Rel. 36, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
15-JUL-1998 (Rel. 36, Last annotation update)
15-JUL-1998 (Rel. 36, Created)
15-JU
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Eukaryota; Metazoa; Nemat
Rhabditidae; Peloderinae;
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Hypothetical protein.
SEQUENCE 371 AA; 41906 MW;
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01-NOV-1997 (Rel. 35, Last sequence update)
01-NOV-1997 (Rel. 35, Last annotation update)
HYPOTHETICAL 41.9 KDA PROTEIN COSD10.4 IN CHROMOSOME
Blattner F.R., Plunkett G.
                         SEQUENCE FROM N.A.
STRAIN-K12 / MG1655;
MEDLINE-97426617; PubMed-9278503;
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                                                                                                                                                                                                                                           Bacteria; Proteobacteria; gamma subdivision; Enterobacteriaceae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Submitted (AUG-1994) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A. STRAIN-BRISTOL N2;
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                                                                                                                                                                   NCBI_TaxID=562;
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Perna N.T.,
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                                                                                               Human adenovirus type 9. Viruses; dsDNA viruses,
                                                                                                                                                                                                                                                                                                                             TRANSMEM SEQUENCE
           viral early region 4 gene.";
Breast Cancer Res. Treat. 39:57-67(1996)
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TRANSMEM
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                                            Javier R.,
                                                      MEDLINE=96327849; PubMed=8738606
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Mizobuchi
                                                                 SEQUENCE FROM N.A.
                                                                                       NCBI_TaxID=10527;
                                                                                                                                                                 P89083;
                                                                                                                                                                                                                                                                                                                                                                                             Hypothetical TRANSMEM
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                                "Mammary tumors induced by human adenovirus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sivasundaram S., Tagami H., Takeda J., Takemoto K., Wada C., Yamamoto Y., Horiuchi T.;
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Gregor J., Davis N.W., Kirkpatrick
                                                                                                                                                                                                                                                                                                                                                                                                                                        EcoGene;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Science
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   SIMILARITY:
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SUBCELLULAR LOCATION: INTEGRAL MEMBRANE PROTEIN (POTENTIAL).
SIMILARITY: BELONGS TO THE RHT FAMILY.
                                                                                                                                                                                                                                                     YMGSK----TLRGRN 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 and this statement is not removed. Usage by and for commercial requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             the Swiss Institute of Bioinformatics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         non-profit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kimura S., Kitakawa M., Kitagawa M., Makino K., Miki T. K., Mori H., Mori T., Motomura K., Nakade S., Nakamura H., Nishio Y., Oshina T., Saito N., Sampei G., Seki Y.,
                                           Shenk T.;
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212 AA;
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Hydrolase.
ACT_SITE
SEQUENCE
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the European Bioinformatics Institute. There
use by non-profit institutions as long as a
modified and this statement is not removed. Us
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Hum. Mol. Genet. 7:1703-1712(1998).
-!- FUNCTION: DUAL-SPECIFICITY PHOSPHATASE THAT
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Q9Z2C5;
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PROSITE; PS50056; TYR_PHOSPHATASE_2; 1:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa;
Mammalia; Eutheria;
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HUMAN
                                                                                                                                                                                                                                                                                                                    SEQUENCE OF 1551-1538 FROM N.A.
MEDLINE=94215165; PubMed=6162575;
GU Y. Alder H., Nakamura T., Schichman S.A., Präsad
Saito.H., Croce C.M., Canaani E.;
"Sequence analysis of the breakpoint cluster region
involved in acute leukenia.";
Cancer Res. 54:2326-2330(1994).
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MEDLINE=93046667; PubMed=1423624;
MEDLINE=93046667; PubMed=1423624;
Tkachuk D.C., Kohler S., Cleary M.L.;
"Involvement of a homolog of Drosophila trithorax by 11q23
chromosomal translocations in acute leukemias.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HRX_HUMAN STANDARD; PRT; 3969 AA; Q03164; Q14845; Q16364; Q13743; Q13744; Q9UMA3; Q1-0CT-1993 (Rel. 27, Created) 01-NOV-1995 (Rel. 32, Last sequence update) 01-OCT-2000 (Rel. 40, Last annotation update) 2INC FINGER PROTEIN HRX (ALL-1) (TRITHORAX-LIKE
"The human MIL gene: nucleotide sequence, homology to the Drosophila trx zinc-finger domain, and alternative splicing."; DNA Cell Biol. 14:475-483(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Djabali M., Selleri L., Parry P., Bowe "A trithorax-like gene is interrupted translocations in acute leukaemias.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens (Human)
Eukaryota; Metazoa; (
                                                                              MEDLINE=95315013; PubMed=7794749;
Marschalek R., Greil J., Lochner
                                                                                                                                                                                                                                                            SEQUENCE_OF 1251-1654 FROM N.A. (ISOFORM 14P-18B)
MEDLINE-95322025; PubMed-7598802;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A. MEDLINE=96290553; PubMed=8703835;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MLL OR HRX OR ALL1 OR TRX1 OR HTRX.
                                                                                                                                                                                                                       Rowley J.D.,
                                                                                                                                                                                                                                              Mbangkollo D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=93265134; PubMed=1303259;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE OF 1-1909 FROM N.A. WEDLINE=93390935; PubMed=8378076;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cell 71:691-700(1992).
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
                                                                                                                      SEQUENCE OF 1212-1603 FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tamamoto K., Seto M., Komatsu H., Iida S., Akao Y., Kojima S., Kodera Y., Nakazawa S., Ariyoshi Y., Takahashi T., Ueda R.;
Two distinct portions of LTG19/ENL at 19p13 are involved in t(11;19)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Exon/intron structure of the human ALL-1 (MIL) gene involved in ranslocations to chromosomal region 11q23 and acute leukaemias.";
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1. 93:966-972(1996).
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                                                                        Lochner K.,
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                                                                            Nilson
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        -i- TISSUE SPECIFICITY: HEART, LUNG, BRAIN AND T AND B LYMPHOCYTES,
-i- DISEASE: INVOLVED IN ACCITE LEUKEMIAS BY CHROMOSOMAL TRANSLOCATIONS
-i T(11;19) (Q23;P13.3) THAT INVOLVES MLL AND MLLT1/ENL;
-i (4;11) (Q21;Q23) THAT INVOLVES MLL AND MLLT2/AF4; T(9;11) (P22;Q23)
-i THAT INVOLVES MLL AND MLLT3/AF9; T(6;11) (Q27;Q23) THAT INVOLVES
-i MLL AND MLLT4/AF6; T(11;17) (Q23;Q21) THAT INVOLVES MLL AND
-i MLLT6/AF17; T(x;11) (Q13;Q23) THAT INVOLVES MLL AND MLLT7/AFX1;
-i T(10;11) (Q21;Q23) THAT INVOLVES MLL AND MLLT1/AF10;
-i T(1;11) (Q21;Q23) THAT INVOLVES MLL AND AF10; T(11;19) (Q23;P13.3)
-i T(1;11) (Q21;Q23) THAT INVOLVES MLL AND AF10; T(11;19) (Q23;P13.3)
-i T(1;11) (Q21;Q23) THAT INVOLVES MLL AND AF10; T(11;19) (Q23;P13.3)
-i T(1;11) (Q21;Q23) THAT INVOLVES MLL AND AF10; T(11;19) (Q23;P13.3)
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-i- FUNCTION: POSSIBLY ACTS AS A TRANSCRIPTIONAL REG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CHROMOSOMAL TRANSLOCATION WITH GAS7.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Forster A., Rabbitts T.H.;
"A method for identifying genes within yeast artificial chromosom application to isolation of MLL fusion cDNAs from acute leukaemia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE OF 1421-1540 FROM N.A. MEDLINE-94020842; PubMed-8414518;
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SIMILARITY: CONTAINS 1
SIMILARITY: CONTAINS 3
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ms T.M., Lange B.J., Felix C.A.;
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D14540;
L01986;
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X83604;
S66432;
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8; Conserv
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IPR002857; -
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Q -> E (IN REF. 2).
R -> G (IN REF. 5).
S -> SGTE (IN REF. 6).
S -> C (IN REF. 6).
Q -> H (IN REF. 4).
P -> S (IN REF. 4).
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VRIIPSSKRTDATIAKQLLQRA -> VSSLILNWKSPRKSG
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01-FEB-1996 (Rel. 3
01-FEB-1996 (Rel. 3
01-CCT-2000 (Rel. 4
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Q02151;
Q1-JUL-1993
Q1-JUL-1993
Q1-JUL-1993
Q1-JUL-1998
          "Cloning and sequencing analysis of a full-length protein alpha subunit, SGA1, from soybean."; Plant Physiol. 108:1315-1316(1995).
                                                                                                                                                             Glycine max (Soybean).
Eukaryota; Viridiplantae;
                                                                                                                                                                                       GUANINE NUCLEOTIDE-BINDING GPA1 OR GA1.
                                                                                                                                                                                                                                                                 SOYBN
GBA1_SOYBN
                                                              Kim W.Y.,
                                                                         STRAIN=CV. WILLIAMS;
MEDLINE=95357428; Pu
                                                                                                   SEQUENCE FROM N.A
                                                                                                                           NCBI_TaxID=3847;
                                                                                                                                      Magnoliophyta; eudicotyledons; core eudicots; Rosidae; Fabales; Fabaceae; Papilionoideae; Glycine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pfam; PF00005; ABC_tran; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       use by non-profit institutions as long modified and this statement is not removed: entities requires a license agreement (See or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRAIN=NCDO 2118;
MEDLINE=93015710; PubMed=1400210;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-JUL-1993 (Rel. 26, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
15-DEC-1998 (Rel. 37, East annotation update)
HYPOTHETICAL ABC TRANSPORTER ATP-BINDING PROTEIN IN LEUD 3'REGION.
Lactococcus lactis (subsp. lactis) (Streptococcus lactis).
Bacteria; Firmicutes; Bacillus/Clostridium group; Streptococcaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EMBL; U92974; AAB81917.1; -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               subsp. lactis."
                                                 Lee S.Y.;
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29 MGAKSLIGRN 138
 FUNCTION: GUANINE
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7; Conserv
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ical protein; ATP-binding; Transport.
ical protein; ATP-binding; Transport.
36 43 ATP (BY SIMILARITY).
259 AA; 29713 MW; 49EED150F6A15169 CRC64;
                                                            Cheong N.E.,
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                                                          PubMed=7630957;
N.E., Lee D.C.,
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37,
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NUCLEOTIDE-BINDING PROTEINS (G PROTEINS) ARE
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PROTEIN ALPHA-1 SUBUNIT (GP-ALPHA-1).
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Hayward D.C., Delaney S.J., Campbell H.D., Ghysen A., Benzer S.,
Kasprzak A.B., Cotsell J.N., Young I.G., Miklos G.L.G.;
"The sluggish-A gene of Drosophila melanogaster is expressed in the
nervous system and encodes proline oxidase, a mitochondrial enzyme
involved in glutamate biosynthesis.";
Proc. Natl. Acad. Sci. U.S.A. 90:2979-2983(1993).
                                                                                                                                                                                                                                                                                                                                   004499; 061349;
01-0CT-1993 (Rel. 27, Created)
01-0CT-1993 (Rel. 27, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
PROLINE OXIDASE, MITOCHONDRIAL PRECURSOR (EC 1.5.3.-)
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       MEDLINE-90188272; PubMed-9520435;
Maleszka R., de Couet H.G., Miklos G.L.G.;
"Data transferability from model organisms to hume from functional genomics of the flightless region Proc. Natl. Acad. Sci. U.S.A. 95:3731-3736(1998).
-1- FUNCTION: CONVERTS PROLINE TO DELTA-1-PERGLIJ
                                                                                                                                                                                                                 STRAIN=CANTON-S;
                                                                                                                                                                                                                                                                            Pterygota; Neoptera; Endopterygota; Diptera;
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                                                                                                  SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                  Ephydroidea;
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                                                                                                                                                                                                                                                                                                     Drosophila melanogaster (Fruit fly)
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SUBUNIT: G PROTEINS ARE C
SUBUNIT: G PROTEINS ARE C
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PR01242; GPROTEINAPLT
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ADP-RIBOSYL[1] (BY ACTION C
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Pred. No.
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a; Brachycera;
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                                   to human beings: insights region of Drosophila.";
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ROLINE-5-CARBOXYLATE. SLUGGISH BEHAVIOR.
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                                   This SWISS-PROT entry is copyright. It is produced through a between the Swiss Institute of Bioinformatics and the EMBI the European Bioinformatics Institute. There are no restrict use by non-profit institutions as long as its content is modified and this statement is not removed. Usage by and fentities requires a license agreement (See http://www.isb-sib
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE
EMBL; AF044218;
                       or send an
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Query Match
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                                                                                                             Submitted (JAN-1998) to the EMBI -:- FUNCTION: POSSIBLY INVOLVED
                                                                                                                                                                                                                                                                                                                              15-DEC-1998 (Rel. 37, Created)
15-DEC-1998 (Rel. 37, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
NEUTRAL TREHALASE (EC 3.2.1.28) (ALPHA, ALPHA-TREHALASE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL; AF017777; AAC PIR; A47302; A47302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EMBL; L07330; AAA02748.1; EMBL; AF017777; AAC28410.
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                                                                                                                                                                             SEQUENCE FROM N.A.
STRAIN=74-OR23-1A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oxidoreductase; Proline metabolism; Mitochondrion; Transit peptide
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                                                                                                                                                              D'Enfert C., Bonini
                                                                                                                                                                                                                              NCBI_TaxID=5141;
                                                                                                                                                                                                                                               Sordariales;
                                                                                                                                                                                                                                                               Eukaryota;
                                                                                                                                                                                                                                                                                Neurospora crassa.
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                                                                                                                                            erenzi H.F.
                                                                                                                                                                                                                                                                                                                 ALPHA, ALPHA-TREHALOSE GLUCOHYDROLASE)
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TISSUE SPECIFICITY: MOST ABUNDANT IN DEVELOPING NERVOUS
DEVELOPMENTAL STAGE: EXPRESSED IN DEVELOPING EMBRYO AS V
                                 SIMILARITY: BELONGS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SIMILARITY: BELONGS
                                                                          CATALYTIC ACTIVITY: ALPHA, ALPHA-TREHALOSE + H(2)0 = 2 D-GLUCOSE.
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                                                            SUBCELLULAR LOCATION: CYTOPLASMIC (BY SIMILARITY)
                                                                                                                                                                                                                                                               Fungi;
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                                                                                                                                                                                                                                               Neurospora.
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Pred. No.
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                                                                                                                                             This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Exthe European Bioinformatics Institute. There are no restr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Fraser C.M., Casjens S., Huang W.M., Sutton G.G., Clayton R.A., Lathigra R., White O., Ketchum K.A., Dodson R., Hickey E.K., Gwinn M. Dougherty B., Tomb J.-F., Fleischmann R.D., Richardson D., Peterson J., Kerlavage A.R., Quackenbush J., Salzberg S., Hanson M., van Vugt R., Palmer N., Adams M.D., Gocayne J.D., Weidman J., Vantack T., Watthey L., McDonald L., Artiach P., Bowman C., Gartinb M. G., Cotton M.D., Horst K., Roberts K., Hatch B., Smith M. Vonta J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   O59189;

15-DEC-1998 (Rel. 37, Created)

15-DEC-1998 (Rel. 37, Last sequence update)

15-DEC-1998 (Rel. 37, Last annotation update)

TOPOISOMERASE IV SUBUNIT B (EC.5.99.1.-)
                                                                                                                                                                                                                                                                                                                        Ojaimi C., Davidson B.E., Saint-Girons I., Old I.G., "Conservation of gene arrangement and an unusual organization of genes in the linear chromosomes of the Lyme disease spirochaetes Borrelia burgdorferi, B. garnini and B. afzelli.";
                                                                                                                                                                                                                                                                              Borrelia burgdorferi, B. garinii and B. afzelii.",
Microbiology 140:2931-2940(1994).
-i-FUNCTION: TOPOISOMERASE IV IS ESSENTIAL FOR CHROMOSOME
SEGREGATION. IT HAS RELAXATION OF SUPERCOILED DNA ACTIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BORBU
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                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=95111614; PubMed=7812434;
                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE OF 1-83 FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nature 390:580-586(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              burgdorferi.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Smith H.O., Venter J.C.;
"Genomic sequence of a Lyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE=98065943; PubMed=9403685;
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Bacteria; Spirochaetales; Spirochaetaceae; Bor
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SEQUENCE 728 AA; 841
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                                                                                                                                                                                                          PERFORMS THE DECATENATION EVENTS REQUIRED DURING THE OF A CIRCULAR DNA MOLECULE (BY SIMILARITY). SUBUNIT: COMPOSED OF TWO SUBUNITS: PARC AND PARE. SIMILARITY: BELONGS TO TYPE II TOPOISOMERASE FAMILY.
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L32861; AAC41408.1;
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                                                                              non-profit institutions as long as its content and this statement is not removed. Usage by a requires a license agreement (See http://www.isban email to license@isb-sib.ch).
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01-NOV-1995
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SEQUENCE
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                                                                                                                                                                                                                                   entities requires a license agreement (S or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                            the European Bioinformatics Institute. The use by non-profit institutions as long modified and this statement is not removed
                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=93342067; PubMed=8341694;
Somerville C.C., Colwell R.R.;
"Sequence analysis of the beta-N-acetylhexosaminidase
vulnificus: evidence for a common evolutionary origin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pfam; PF00204; DNA_topoisoII; 1. PRINTS; PR00418; TPI2FAMILY. PROSITE; PS00177; TOPOISOMERASE_II;
                                                                                                                                                                                                 HSSP;
                                                                                                                                                                                                            EMBL;
                                                                                                                                                                                                                                               entities requires a
                                                                                                                                                                                                                                                                                                This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A. STRAIN=ATCC 27562;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bacteria; Proteobacteria; NCBI_TaxID=672;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (BETA-N-ACETYLHEXOSAMINIDASE) (CHITOBIASE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BETA-HEXOSAMINIDASE
                                                                                                                   DISULFID
                                                                                                                               Hydrolase;
DISULFID
                                                                                                                                                         PRINTS; PR00738;
                                                                                                                                                                                                                                                                                                                                                                                                                                hexosaminidases.";
Proc. Natl. Acad.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Vibrio vulnificus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            581
                                                                                                                                                      PF00728; Glyco_hydro_20; PF00738; GLHYDRLASE20.
                                                                                                                                                                                  [nterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 FYMGSKTLRGRN
                                                                                                                                                                                                                                                                                                                                       PATHWAY: CONVERSION OF CHITIN TO GLCNAC. SIMILARITY: BELONGS TO FAMILY 20 OF GLYCOSYL HYDROLASES
                                                                                                                                                                                                                                                                                                                                                                 CATALYTIC ACTIVITY: HYDROLYSIS OF TERMINAL NON-REDUCING N-ACETYL-D-HEXOSAMINE RESIDUES IN N-ACETYL-BETA-D-HEXOSAMINIDES.
                                                                                                                                                                                                                                                                                                                                                                                                     C. Natl. Acad. Sci. U.S.A. 90:6751-6755(1993).
FUNCTION: HYDROLYSIS OF TERMINAL, NON-REDUCING N-ACETYL-BETA-D-GLUCOSAMINE RESIDUES IN CHITOBIOSE AND HIGHER AND
                                                                                                                                                                                                                                                                                                                                                                                          IN GLYCOPROTEINS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FYMGQNTPERRN
                                                                                                                                                                                             L04544; AAA27527.1;
Q54468; 1QBB.
 Similarity 6; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity 7; Conserv
                                                                                        ; Glycosidase.

31 40

377 385

484 530

519 519
                                                                                                                                                                                 IPR001540;
                                                                847 AA;
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599 AA; 68774
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (Rel. 32, Created)
(Rel. 32, Last sequence (Rel. 39, Last annotation)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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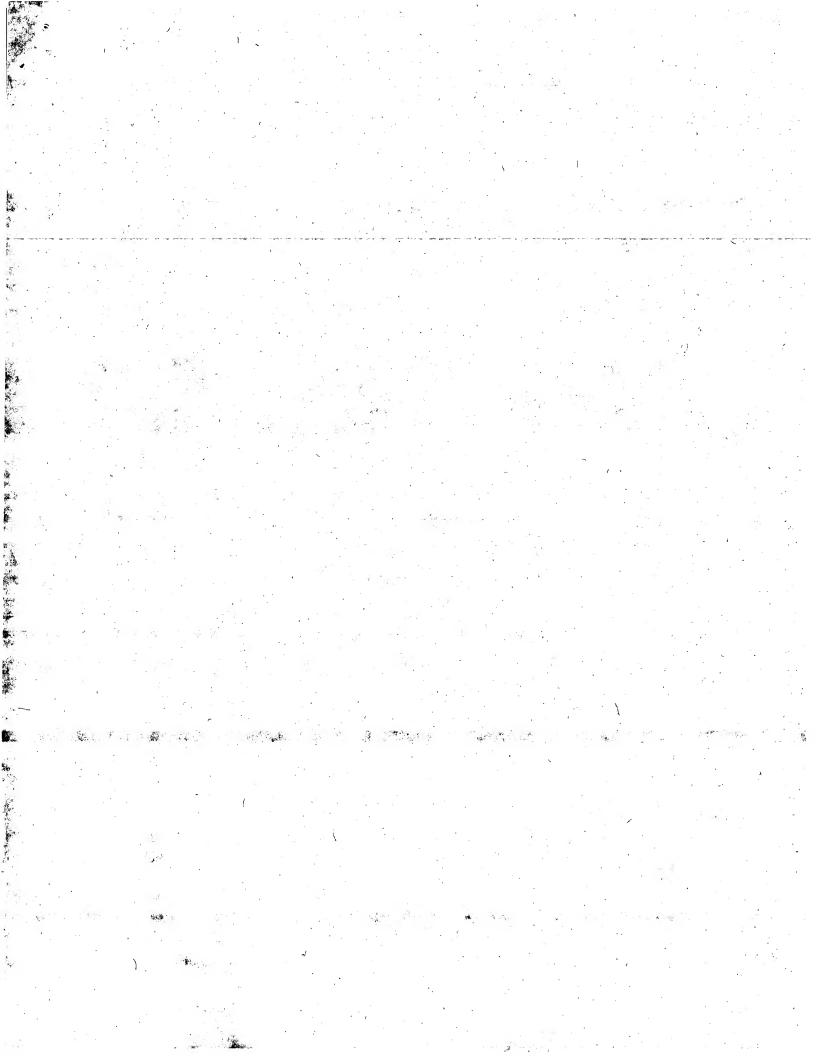
    Last sequence update)
    Last annotation update)
    (EC 3.2.1.52) (N-ACETYL-BETA-GLUCOSAMINIDASE)

                                                                 94277
             54.7%;
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ATP-binding.
MW; B5901F17B1CC7721 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gamma subdivision; Vibrionaceae;
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Pred. No.
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BY SIMILARITY.
CATALYTIC ACID (BINDS TO THE GLYCOSIDIC LINKAGE) (BY SIMILARITY).
3C4405EFA34F3D14 CRC64;
             Score 35; DB Pred. No. 43;
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 Mismatches
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Search completed: June 4, 2001, 12:24:05 Job time: 567 sec



GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.

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Minimum
Maximum
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Perfect score:
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Listing first 45 summaries
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sp_bacteria:*
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sp_vertebrate:*
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13.062 Million cell updates/sec
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Q91nv4 arabidopsis
Q9vrh9 drosophila
Q9vrh8 drosophila
Q9vrh8 drosophila
Q43546 lilium long
Q61818 mus musculu
Q9u1d0 leishmania
Q91uy1 arabidopsis
Q9kxc4 escherichia
                                                                                                      Q26654 sarcophaga
Q13744 homo sapien
Q77335 plasmodium
Q47307 escherichia
                                                                                                                                                                        Q9mzt0
                                                                                                                                                                                                          Q92663 homo sapie
Q92495 homo sapie
O58234 pyrococcus
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                                                                                                                                                                                                                                     homo sapien
                                                                                                                                                                                     arabidopsis
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ALIGNMENTS

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RESULT 2.
Q92495
ID Q92495
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Q92663
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Best Local Similarity
Matches 12; Conserv
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01-FEB-1997
01-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                             Q92663
Q92663;
                                                                                                                                                                                                                                                           MEDLINE-93055454; PubMed=1430234; Porges A.J., Redecha P.B., Doebele R., Kimberly R.P.;
                                                                                                                                                                                                                                                                                                                              Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                 01-OCT-2000 (TrEMBLrel. FC GAMMA RECEPTOR I.
                                                                                                                                                         SEQUENCE
                                                                                                                                                                         INTERPRO; IPRO03006; -. PFAM; PF00047; ig; 3.
                                                                                                                                                                                                   EMBL; L03418; AAA36049.1;
HSSP; P12319; LALT.
                                                                                                                                                                                                                                     cells.";
                                                                                                                                                                                                                                                                                            TISSUE-BLOOD;
                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
                                                                                                                                                                    PRODOM; PD002534;
                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=9606;
                                                      1 FYMGSKTLRGRN 12
|||||||||
229 FYMGSKTLRGRN 240
                                                                                                                                                                                                                          . Clin. Invest. 90:2102-2109(1992)
                                                                                                                                                                                                                                               Novel
                                                                                                                                                                                                                                               Fc gamma receptor I family gene products in human mononuclear
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                                                                                                  Conservative
 PRELIMINARY;
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                                                                                                                                                         42632 MW;
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15,
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Last sequence update)
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                                                                                                             Score 64;
Pred. No.
  PRT;
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                                                                                                    Mismatches
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  375
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                                                                                                             0.00026;
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Matches 12
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01-JUN-2000
Kawarabayasi Y., Sawada M., Horikawa H., Haikawa Y., Hino Y., Yamamoto S., Sekine M., Baba S.-I., Kosugi H., Hosoyama A., Naga Yamamoto S., Ogura K., Otsuka R., Nakazawa H., Takamiya M., Ohfuku Sakai M., Ogura K., Otsuka R., Kudoh Y., Yamazaki J., Kushida N., Oguc Funahashi T., Tanaka T., Kudoh Y., Yamazaki J., Kushida N., Oguc Aoki K.-I., Yoshizawa T., Nakamura Y., Robb F.T., Horikoshi K.,
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058234;
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Ol-FEB-1997 (TrEMBLrel. 02, Last sequence update)
Ol-FCT-2000 (TrEMBLrel. 15, Last sequence update)
FC GAMMA RECEPTOR TYPE I (FC GAMMA RIB-FC GAMMA RECEPTOR)
                                                                                                                                                                                                                                                                                                                                                             PFAM; PF00047; 1g: 3
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01-FEB-1997
                                                                 STRAIN-OT3
                                                                           SEQUENCE
                                                                                               Archaea; Euryarchaeota; NCBI_TaxID=53953;
                                                                                                                   Pyrococcus horikoshii.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=93018827;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE OF 1-224 FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Submitted (DEC-1992) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ezekowitz
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NCBI_TaxID=9606;
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                                                   MEDLINE=98344137; PubMed=9679194;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
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12; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A.B.:
                                                                                                                                      (TTEMBLrel. 07, Created)
(TTEMBLrel. 07, Last sequence up
(TTEMBLrel. 14, Last annotation
L 44.0 KDA PROTEIN PH0498.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ); AAA58414.1;
                                                                                                                                                                                            PRELIMINARY,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                176:1115-1123(1992).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sastry K., Iyer R.R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sastry K.N., Iyer R.R.,
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AAD13842.
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01-AUG-1998
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Q9LPL9;
Q1LPL9;
Q1-OCT-2000 (TrEMBLrel. 15;
Q1-OCT-2000 (TrEMBLrel. 15;
Q1-OCT-2000 (TrEMBLrel. 15;
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Arabidopsis thaliana (Mouse-ear cress).
Eukaryota; Viridiplantae: Embryophyta; Tracheophyta; Spermatophyta;
Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids II;
Brassicales; Brassicaceae; Arabidopsis.
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PROTEIN PH0394.
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horikoshii OT3.";
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RESULT
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O9Z2C5;
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01-MAY-1999 (TrEMBLrel. 1
01-MAY-2000 (TrEMBLrel. 1
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Sakano H., Liu S.X., Yu G., Lee J., Lenz C., Pham P., Toriumi M.,
Chin C., Chlou J., Choi E., Chung M., Gonzalez A., Howng B., Liu A.,
Chin C., Chlou J., Choi E., Chung M., Gonzalez A., Howng B., Liu A.,
Vaysberg M., Altafi H., Brooks S., Buehler E., Chao Q., Conn L.,
Conway A.B.; Hansen N.F., Johnson-Hopson C., Khan S., Kim C., Lam B.,
Miranda M., Nguyen M., Palm C., Shinn P., Southwick A., Davis R.W.,
Ecker J.R., Federspiel N.A., Theologis A.;
"The sequence of BAC F2210 from Arabidopsis thaliana chromosome 1.";
Submitted (JUN-2000) to the EMBL/GenBank/DDBJ databases.
EMBL; AC015445; AAF76448.1;
EMBL; AC015445; AAF76448.1;
                                                                                                                                                                                                                                                                                                                                                                        Yan Y., Li X., Wang A., Zhang G.;
"Molecular cloning and identification of full-length cDNA encoding high affinity Fc receptor for bovine IgG (Fc gamma RI).";
Vet. Immunol. Immunopathol. 75:151-159(2000).
EMBL, AF162866; AAF80477.1; -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bos taurus (Bovine).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
Bovidae; Bovinae; Bos.
                                     Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                               Mus musculus (Mouse).
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SEQUENCE FROM N.A
                         NCBI_TaxID=10090;
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                                     Chordata;
Rodentia;
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87.5%;
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                                    Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                             PRT;
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RESULT Q13744
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Q13744; Q13743;
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INTERPRO;
                       Homo sapiens (Human).
Eukaryota; Metazoa; C
Mammalia; Eutheria; F
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01-NOV-1996 (TrEMBLrel.
01-NOV-1996 (TrEMBLrel.
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Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;
Pterygota; Neoptera; Endopterygota; Diptera; Brachycera; Musc
Oestroidea; Sarcophagidae; Sarcophaga.
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SEQUENCE
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                                                                                           ALL-1.
                                                                                                                ALL-1 PROTEIN.
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SEQUENCE 116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             translational processing mechanism.";
J. Biol. Chem. 270:4624-4631(1995).
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Mol. Genet. 7:1703-1712(1998).
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P04253; 10XY.
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"Molecular analysis of the chromosomal breakpoint and fusion
"transcripts in the acute lymphoblastic SEM cell line with chromosomal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Marschalek R., Greil J.,
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Similarity 66. 8; Conservative
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PS00354; HMGI_Y; UNKNOWN_1
PS50014; BROMODOMAIN_2; 1.
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                57.8%;
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                DB 4; 5.6e+02;
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                               Length 4005;
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Mismatches

Indels

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Gaps

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2 YMGSKTLRGRN

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RESULT
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Best Local S
Matches 6
Query Match
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Matches 7
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01-NOV-1996
01-MAY-1999
01-MAY-1999
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01-MAY-1999
01-MAY-2000
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                                                      origins for three different submitted (AUG-1998) to the EMBL; AJ008006; CAA07835.1; Plasmid.
                                                                                                                                                                                                                                                                                                                                                                                                                      Mungall K., Lawson D., Barrell B.;
Submitted (SEP-1998) to the EMBL/GenBank/DDBJ da
EMBL; 298547; CAB11120.2; -
INTERPRO; IPR001064; -
PROSITE; PS00225; CRYSTALLIN_BETAGAMMA; UNKNOWN_
SEQUENCE 280 AA; 33089 MW; 1C37A22BCD82A846
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Plasmodium falciparum
Eukaryota; Alveolata;
NCBI_TaxID=5833;
                                              SEQUENCE
                                                                                                                                                         TRANSPOSON=IS1294;
                                                                                                                                                                                                               Escherichia coli.
Plasmid pUB2380.
                                                                                                                                                                                                                                   TNP1294
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PFC0410W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PFC0410W PROTEIN.
                                                                       Albiger B., Bennett P.M., Commanducci A., Dodd H.M., Lett M.C.; "pUB2380 : a ColD-like resitance plasmid with three replication origins for three different functions."; Submitted (AUG-1998) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                            TRANSPOSASE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
                                                                                                                                         Submitted
                                                                                                                                                                                                      Bacteria; Proteobacteria;
                                                                                                                                                   Bruyns E.;
                                                                                                                                                                                     NCBI_TaxID=562;
                                                                                                                                                                                                Escherichia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             872
                                                                                                             RANSPOSON=IS1294;
Local Similarity 63. hes 7; Conservative
                                                                                                                                                                                                                                                                                                                                      98 FYYDNKVFRGKN 109
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                                               351 AA;
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                                                                                                                                                                                                                                                                                          PRELIMINARY;
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                                               40283 MW;
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                                                                                                                                                                                                        gamma subdivision; Enterobacteriaceae;
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Last sequence update)
Last annotation update)
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Score 36; DB Pred. No. 64; 0; Mismatches
  0;
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Pred. No.
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                                              46FAD7C01E075CAD CRC64;
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                    Length 351;
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  Indels
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237

YMSKKTAGGRN

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Query Match
Best Local Similarity
"--hes 6; Conserv
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01-MAY-2000
01-JUN-2000
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Q9LNV4;
01-OCT-2000
                                                                                                  Q9VRH9;
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                                                                                                                                                                                                                                                                                                                                                                                                                          Cheuk R., Shinn P., Brooks S., Buehler E., Chao Q., Johnson-Hopson C., Khan S., Kim C., Altafi H., Bei B., Chin C., Chiou J., Choi E., Conn L., Conway A., Gonzalez A., Hansen N., Howing B., Koo T., Lam B., Lee J., Lenz C., Li J., Liu A., Liu J., Liu S., Mukharsky N., Nguyen M., Palm C., Pham P., Sakano H., Schwartz J., Southwick A., Thaveri A., Toriumi M., Vaysberg M., Yu G., Davis R., Federspiel N.,
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Drosophila melanogaster (Fruit fly)
                                   SLGA PROTEIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Viridiplantae; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids II;
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Ecker J.R.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Submitted
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Submitted (FEB-2000) to the EMBL/GenBank/DDBJ databases.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted (MAR-2000)
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                                                                                                                                                                                                                                     2 YMGSKTLRGRN 12
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Pred.
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196
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                                                                                                                  669' AA.
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                                                     Q9VRH8;
01-MAY-2000 ()
01-MAY-2000 ()
01-JUN-2000 ()
SLGA PROTEIN.
 Drosophila ı
Eukaryota; ı
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S. Berkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P., Burtis K.C., Busam D.A., Butler H., Cadieu E., Center A., Chance Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P., de Pablos B., Delcher A., Deng Z., Mays A.D., Dew I., Dietz S., Dodson K., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Merkulov G., Milshina N.V., Mobarry C., Morris J., Moshrefi A., Mount S.M., Moy M., Murphy B., Murphy L., Muzny D.M., Nelson D. Nelson D. R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M., Palazzolo M., Pittman G.S., Pan S., Pollard J., Puri V., Reese J. Reinert K., Remington K., Saunders R.D.C., Scheeler F., Shen H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hostin D., Houston K.A., Howland T.J., Wei M.-H., Ibegwam C.,
Hostin M., Kalush F., Karpen G.H., Ke Z., Kennison J.A., Ketchum
Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z.,
Lasko P., Lei Y., Levitsky A.A., Li J., Li Z., Liang Y., Lin X.,
Liu X., Mattel B., McIntosh T.C., McLeod M.P., McPherson D.,
Liu X., Mattel B., McIntosh T.C., McLeod M.P., McPherson D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Durbin K.J., Evangelista C.C., Ferraz C., Ferriera S., Fleischmann Fosler C., Gabrielian A.E., Garg N.S., Gelbart W.M., Glasser K., Glodek A., Gong F., Gorrell J.H., Gu Z., Guan P., Harris M., Harris N.L., Harvey D., Heiman T.J., Hernandez J.R., Houck J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Amanatides P.G., Scherer S.E., Li P.W., Hoskins R.A. George R.A., Lewis S.E., Richards S., Ashburner M., Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Brandon R.C., Rogers Y.-H.C., Blazej R.G., Champe M.
                                                                                                                                                  Q9VRH8
                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Science
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Beeson K.Y., Benos P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Brandon R.G., Rogers Y.-H.C., Blazej R.G., Champe M., Prelitet B.J., Wan K.H., Doyle C., Baxter E.G., Helt G., Nelson C.R., Miklos G.L.G., Wan K.H., Doyle C., Baxter E.G., Helt G., Nelson C.R., Miklos G.L.G., Wan K.H., Doyle C., Baldwin D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Pterygota; Neoptera; Endopterygota; Diptera; Brachycon Ephydroidea; Drosophilidae; Drosophila.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FLYBASE; FBgn0003423; slgA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "The genome sequence of Drosophila Science 287:2185-2195(2000). EMBL; AE003568; AAF50819:1; -.
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FYFGAKLVRG 500
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Similarity 60.0%;
6; Conservative
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 melanogaster (Fruit
Metazoa; Arthropoda;
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AA; 75951 MW;
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V., Berman B.P., Bhandari D., Bo
                                                                         13,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Yandell M.D., Zhang Q., Chen L.X., C., Blazej R.G., Champe M., Pfeiffer B.D.
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                                                                                                                                                  PRT;
fly).
Tracheata; Hexapoda;
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     Insecta;
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asley E.M.,
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LIMY.
LIMY Iongiflorum (Trumpet lily).
Lilium longiflorum (Trumpet lily).
Eukaryota; Viridiplantae; Embryophyta; Tracheophyta; Spermatophyta;
Eukaryota; Liliopsida; Liliales; Liliaceae; Lilium.
Magnoliophyta; Liliopsida; Liliales;
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Hostin D., Houston K.A., Howland T.J., Wel M.-H., Ibegwam C.,
Jalali M., Kalush F., Karpen G.H., Ke Z., Kennison J.A., Ketchum K.A.,
Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z.,
Liako P., Lei Y., Levitsky A.A., Li J., Li Z., Liang Y., Lin X.,
Liu X., Mattel B., McIntosh T.C., McLeod M.P., McPherson D.,
McIntu X., Mattel B., McIntosh T.C., McLeod M.P., McPherson D.,
Merkulov G., Milshina N.V., Mobarry C., Morris J., Moshrefi A.,
Mount S.M., Moy M., Murphy B., Murphy L., Muzny D.M., Nelson D.L.,
Nelson D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M.,
Nelson D.R., Nelson K., Saunders R.D.C., Scheeler F., Shen H.,
Shine R.C., Siden Riamost T., Saunders R.D.C., Scheeler F., Shen H.,
Shine R.C., Siden Riamost T., Saunders R.D.C., Scheeler F., Shen H.,
Shine R.C., Siden Riamost T., Saunders R.D.C., Scheeler F., Shen H.,
Shine R.C., Siden Riamost T., Saunders R.D.C., Scheeler F., Shen H.,
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Shine R.C., Siden Riamost T., Shen H.,
Shine R.C., Siden Riamost T., Shen H.,
Shine R.C., Siden R.J., Shen H.,
Shine R.C., Shen H.,
                                                                                                                                                                                        SERINE PROTEINASE (FRAGMENT).
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INTERPRO; IPR002872; -
PFAM; PF01619; Pro_dh; 1.
SEQUENCE 671 AA; 76113 MW;
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"The genome sequence of Drosophila melanogaster.";
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Sutton G.G.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ETTERCE 28/:2185-2195(2000).
EMBL; AE003568; AAF50820.1;
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Amanatides P.G.,
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Ephydroidea; Drosophilidae; Drosophila.
NCBI_TaxID=7227;
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Baxter E.G., Helt G., Nelson C.R., Miklos G T O.

A. An H.-U., Andrews-Dfn...
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Pred. No. 1.3e+02;
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wiklos G.L.G.
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MENDEL; 11876; Lillo;1086;11876.
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                                                                                                                                                                                                                                                                                            microsporocytes.
                                                                                                                                                                                                                                                                                                                                        STRAIN-CV. HINOMOTO; TISSUE-FLORAL BUD; MEDLINE-96051386; PubMed=7584025;
                                                                                                                                                                                                                                                                                                                                                                                                           STRAIN=CV. HINOMOTO; TISSUE=FLORAL BUD;
                                                                                                                                                                            PFAM; PF00082; Peptidase_S8; 2.
PRINTS; PR00723; SUBTILISIN.
PROSITE; PS00136; SUBTILASE_ASP;
PROSITE; PS00138; SUBTILASE_SER;
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                                                       YYSGHTLRGRN 515
                                                                             YMGSKTLRGRN 12
                                                                                                  Similarity 63.07; Conservative
                                                                                                                                                                                                                                                                                                                                                                                   (DEC-1993) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                          813 AA;
           June
                                                                                                                                                                                                                                                                                                       of cDNAs induced in meiotic prophase in lily
           4,
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          2001, 12:23:26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     6.463 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              390729
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  83.DAT:
                                                                             Oligopeptide from Human sFC-gammaRI Human macrophage-s Human FCRI (CDNA c Human macrophage-s Human macrophage-s
                                                                                                                                                                                                      Description
              Human
                                Human
                                                Human
                                                               Human
                            n macrophage-s
n FcRI (cDNA c
n FcRI (cDNA c
              receptor
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	31	31	31	32	32	32	32	32	32	32	32	32	32	32	3	3	33	33	33	33	ω ω	ω S	34	36	36	φ	36	36	36	57	57	57	57	57
	54.4	54.4	54.4	56:1	56.1					56.1			56.1	5	7.	57.9	7	57.9	7	.7	57.9	7	٥	ū	ω ·		Ü	Ü	٠	•		100.0	٠	100.0
	255	185	58	992	969	, 802	623	525	506	451	428	284	283	25	1165	749	696	696	395	371	35 35	193	88	2004	2004	2004	2004	1879	410	399	377	374	374	374
	21	21	21	21	21	21	15	20	21	21	18	21																14	12	21	20	21	21	19
	Y74710	Y58587	G00321	G33920	G33921	G33922	R49578	Y25881	Y94986	G39206	W11883	G39207	G39208	Y25887	R37309	P70286	Y17207	W55698	Y31732	B54247	Y67867	W55378	Y35386	B10469	W89402	W37052	R99464	R42455	R12428	B43683	W86195	Y96226	Y96134	W97834
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			,	:	•	٠																												
	Neisseria		Human	Arabidopsis	Arabidopsis	Arabidopsis	Sequence of	Human	. Human secre	Arabidopsis	Hantavirus	Arabidopsis	Arabidopsis	Human		Protein en						H. pylori				S. putrefaci					Human	Human	Human	Human

ALIGNMENTS

RESULT W60557 W60557; W60557 standard; peptide; 12 AA. 07-OCT-1997; W09815833-A1. Synthetic. immunohistochemical; immunoflu
cell transformation; mutation; Oligopeptide from extracellular domain of CD64 18-AUG-1998 De Kruif CA, (UYUT-) RIJKSUNIV UTRECHT. 08-OCT-1996; Extracellular domain; 16-APR-1998 (first entry) Logtenberg 96EP-0202791 97WO-NL00557 immunofluorescent analysis; detection; CD64; identification; antibody; oncogene.

Human

receptor

Identifying peptide(s) binding specifically to protein target - by expressing on phage surface and testing for binding to immobilised oligo:peptide derived from the target, useful for, e.g. identifying specific antibodies

WPI; 1998-240964/21.

method for the identification of a peptide able to bind specifically to a target protein. The method comprises displaying the peptide on the surface of a replicable display package, synthesising oligopeptides derived from the target protein on a solid phase, and testing for binding between the peptide and oligopeptides. The method is used to screen_large peptide libraries, especially to detect antibodies, or their fragments, that bind to cell markers or that can differentiate between different forms of the same protein, including bispecific antibodies that bind to wo non-overlapping epitopes on the same monomeric antigen or two epitopes on different molecules. The genes/oligonucleotides that encode selected peptides can be isolated and used for recombinant production of antibodies. Selected antibodies are used, e.g. for immunohistochemical or immunofluorescent analysis, and also to detect cell transformation caused by mutation in 'antioncoenes.'

domain of CD64. They were synthesised on 25 polyethylene rods adjacent, non-overlapping sets. Peptide specific phages were affinity binding to the CD64-covered rods. The specification

Synthetic oligopeptides W60537-61 are derived from the extracellular

25 polyethylene rods as

The specification describes

isolated

by s a

This invention describes a novel model of an Fc receptor (FCR) protein representing a three-dimensional (3-D) structure that substantially

conforms to the specified atomic coordinates. Computer model images of

Claim 11; Page 316-317; 326pp; English

Example 1; Page 29; 40pp; English

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ΧX
WPI; 1999-539978/45
                                                       Maxwell KF,
                                                                                                                                              11-SEP-1998;
06-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                        atomic coordinate; bioactive compound design; computer-assisted; drug design; therapy; inhibitor; Fc-gammaR; Fc-epsilonR; IgG; IgE; tissue damage; hypersensitivity; inflammatory cell recruitment;
                                    Powell MS
                                                                      Baell JB,
                                                                                                        (ILEX-) ILEXUS PTY LTD.
                                                                                                                                                                                                   04-FEB-1999;
                                                                                                                                                                                                                                                                         W09940117-A1
                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                               anti-inflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Fc receptor; Fc-gammaRIIa; human; FCR; model; three-dimension;
                                                                                                                                                                                                                                     12-AUG-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human sFc-gammaRI protein fragment.
                                                                                                                                                                                                                                                                                                                                                                  inflammatory
                                                 Epa V, Garrett, McCarthy TD,
                                                                                                                                                                                                                                                                                                                                                                  modulator; Fc-gammaRIa;
                                                                                                                                            98US-0099994.
98US-0073972.
                                                                                                                                                                                                 99WO-IB00367.
                                                                                                                                                                                                                                                                                                                                               immunoprotective;
                                                                                                                                                                                                                                                                                                                                                                                                                bioactive compound design;
                                                 TPJ, Hogarth PM,
McKenzie IFC, Pi
                                                                                                                                                                                                                                                                                                                                            immune function sFc-gammaRI.
                                               Pietersz
                                                                    Matthews
                                                                                                                                                                                                                                                                                                                                                               regulation;
                                                                    BR;
                                                                                                                                                                                                                                                                                                                                                                                                                                 3-D;
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Three-dimensional structures and models of Fc receptors, useful in computer-assisted drug design

This amino acid sequence was predicted from the cDNA sequence of cDNA clone p98/X2. It differs from the sequence predicted from clone p135 (see Q21178) at position 25, p135 encodes a Ser residue while clone p98/X2 predicts a Thr residue. At position 58, p90

Example 10; Page 94a; 160pp; English

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RESULT
R22549
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Best Local
                                                                                                         New CD53 cell surface antigen and DNA encoding it - for immuno-therapy and diagnosis of haematopoietic neoplasms,
                                                                                                                                                            N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     reduce IgG- or IgE-mediated tissue damage, e.g. IgG- or IgE-mediated hypersensitivity, recruitment of inflammatory cells or release of inflammatory modulators. The therapeutic compositions can also be used to treat disease in animals. The knowledge of the 3-D structure of Fc-gamma RIA and models of other FcR provides a means for designing and producing compounds that regulate immune function and inflammation in an animal, including humans (i.e. structure based drug design). For example, chemical compounds can be designed to block binding of immunoglobulin to chemical compounds can be designed to block binding of immunoglobulin to
                                                                                                                                                            WPI; 1992-056864/07
N-PSDB; Q21179.
                                                                                                                                                                                                           Seed B,
                                                                                                                                                                                                                                                                           13-JUL-1990;
                                                                                                                                                                                                                                                                                                          15-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                         WO9201049-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human macrophage-specific FcRI receptor encoded by clone p98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  R22549 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          an PC receptor protein using various computer programs and models. The products of the invention have anti-inflammatory and immunoprotective activity. This sequence represents the human sFC-gammaRI protein fragment described in the method of the invention.
                                                                                                                                                                                                                                           (GEHO-) GEN HOSPITAL CORP.
                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                      Rapid immunoselection cloning technique; cell surface antigen; immunodiagnosis; high affinity receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     oligonucleotides, peptides, peptidomimetics and small organic moleculoby computer-assisted methods of drug design. Therapeutic compositions that inhibit the activity of Fc-gammaR or Fc-epsilonR can be used to reduce 196- or IgE-mediated tissue damage, e.g. 196- or IgE-mediated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  the FcR can be used to design bioactive chemical compounds,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 TSSEYQILTARR 12
|||||||||
222 tsseyqiltarr 233
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 l Similarity
12; Conser
                                                                                                                                                                                                           Aruffo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               261
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                           90US-0553759
                                                                                                                                                                                                                                                                                                        90WO-US04986
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                                                                                                                                                                                                             Amiot M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    344
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 57; DB 20; I
Pred. No. 0.00078;
Pred. No. 0.00078;
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1 TSSEYQILTARR 12

15-NOV-1999 Y33183

(first entry)

Y33183 standard;

Protein; 261 AA

N

Matches Query Match

Local Similarity 100 nes 12; Conservative

100.0%;

Score 57; DB 19; Pred. No. 2.6e-05; Mismatches

Length 12; Indels

0;

0;

0;

Gaps

Sequence

12

by mutation in (anti)oncogenes.

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RESULT
R91439
ID RS
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    Query Match
Best Local 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (see Q21180) predicts Val and p98/X2 predicts Leu. Sequences predicted from all 3 clones show the typical features of a type I integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane-spanning domain, and a short, highly charged cytoplasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and six Cys residues distributed among three C2 set Ig-related domains. A fusion protein of FCRI and a receptor ligand will be helpful to
            The amino acid sequence (R91439) of human FCRI was detd. from a CDNA library using an immunoselection cloning method. FCRI is a high affinity receptor for the Fc portion of IgG, normally located on cell surfaces of macrophages. 2 Other CDNA clones, pl35 (T14717) and p90 (T14719), coded for a variants of the FCRI sequence; the C-terminal sequence
           macrophages. 2 Other cDNA clones coded for a variants of the FcRI
                                                                                                                                     Cloning of cDNA encoding cell surface of diagnostic and therapeutic proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                       Aruffo
                                                                                                                                                                                                                                                                            13-JUL-1990;
18-OCT-1993;
                                                                                                                                                                                                                                                                                                                                  01-DEC-1992;
                                                                                                                                                                                                                                                                                                                                                              25-FEB-1988;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cell surface antigen; cloning; immunoselection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human FcRI (cDNA clone p98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R91439;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R91439 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          increase the potency of antibodies
                                                                                                          Example 10; Column 55-56; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                   US5506126-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             therapy;
                                                                                                                                                                                                                                                 (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                       L3-JUL-1989
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les 12; Conserv
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DB; T14718.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           diagnosis;
product
                                                                                                                                                                                                                       Seed
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                      92US-0983647.
88US-0160416.
89US-0379076.
                                                                                                                                                                                                                                                                            900S-
                                                                                                                                                                                                                                                                                                                                                              88US-0160416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Protein;
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                                                                                                                                                                                                                                                                            s-0553759.
s-0139273.
 is truncated compared
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             vector;
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                                                                                                                                                                                                                                                                                                                                                                                                                                             "amino acid 58 is Leu in 
'translated product"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    product).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FcRI; Fc receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ₿
                                                                                                                                                   surface antigen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           in therapy.
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 with
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          clone p135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        immunotherapy;
                                                                                                                                                     useful for isolation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 344;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
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RESULT
Y96183
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FΤ
                                                                                                                                                                                                                                                   77777
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      and p90 products (see also R91438 and W00859). of FCRI and a receptor ligand will be helpful i potency of antibodies in therapy.
Misc-difference
                  Misc-difference
                                                      Misc-difference
                                                                        Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Y96183 standard;
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                                   Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                            panning; immunodiagnosis; diagnosis; immunotherapy; gene th
immune disorder; infection; asthma; immune-complex disease;
                                                                                                                                                                                                                                                                                                                                                                                                                               Macrophage;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-DEC-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ch 100.
1 Similarity 100.
12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       344 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                               FCRI; cell surface antigen; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                   multiple sclerosis; parasitic disease;
                                                                                                                                                                /note=
117
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116
                /note=
140
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                                    139
                                                                                                                     /note=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                     Fc receptor
                                                                                                                                                                                                                                                                                                                                                                                                                      gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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Ъ

241

tsseyqiltarr

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25-FEB-1988;
13-JUL-1989;
23-MAR-1990;
Sequence
                                                                surfaces other than macrophages,
                                                                                                                                                                                                    (panning on) an antibody-coated substrate such as a culture dish. CSA nucleic acids isolated by the method of the invention, and the proteins they encode, are useful for immunodiagnostic and
                                                protein of FcRI and a receptor
                                                                         animals, including humans. These disorders include asthma, immune-complex disease, amyloidosis, parasitic diseases or multiple sclerosis. FCRI is a high affinity receptor for the FC portion of IGG, normally located on the cell surfaces of macrophages. The ability to interfere with such bonding, or to cause it to occur on surfaces of the relation macrophages.
                                                                                                                                                   animals, including humans.
                                                                                                                                                                                                                                                      physical selection of cells expressing the antigen by adhesion to
                                                                                                                                                                                                                                                                          based upon transient expression of a CSA in eukaryotic cells and
                                                                                                                                                                                                                                                                                                                                                                                                                             Isolated nucleic acid molecule encoding the CD19 cell surface antigen, useful for immunodiagnosis and immunotherapy of immune-mediated infections or disorders, e.g. asthma, immune-complex disease, parasiting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2000-586382/55
N-PSDB; A50631.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Stamenkovic I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-DEC-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US6111093-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Misc-difference
                                                                                                                                                                     treatment of
                                                                                                                                                                                                                                                                                                              trom a cDNA library expressed
                                                                                                                                                                                                                                                                                                                                           The present sequence
                                                                                                                                                                                                                                                                                                                                                                               Example 10; Column 53-55; 75pp;
                                                                                                                                                                                                                                                                                                                                                                                                                   diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             28-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-AUG-2000
                                                                                                                                                                 mmunotherapeutic applications, including reatment of immune mediated infections,
                                                                                                                                                                                                                                                                                                                              deduced from
                                potencies
                                                                                                                                                                                                                                                                                          invention designed to isolate CSA nucleic acids.
344 AA;
                                  of.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         90US-0498809
90US-0553759
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           89US-0379076
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           92US-0983647
88US-0160416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             98US-0181612
                                                                                                                                                                                                                                                                                                     uence is that of a human macrophage specific FCRI, cDNA clone p135 (see A50631), which was isolated rary expressed in COS cells using a novel method of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note=
338
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note=
306
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note=
305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note=
268
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220
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216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Seed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      333
                                antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    В;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   encoded by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "encoded by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "encoded by GTG"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "encoded by CGT"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "encoded by TTC"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "encoded by
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "encoded by
                          nages, is useful intor ligand will be in therapy.
                                                                                                                                                                   including the diagnosis and fections, diseases, and disorders in
                                                                                                                                                                                                                                                                                                                                                                               English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CCT"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GGC"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAC "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAT"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CAA"
                                           in therapy. A fusion be helpful to increase
                                                                                                                                                                                                                                                                                          The method is
                                                                                                                                                                                                                                                                                                                                                                                                                               parasitic
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Matches Query Match Best Local S

l Similarity 100 12; Conservative

100.0%;

Score 57; DB 21; Pred. No. 0.0011;

Length 344; Indels ·

0;

Gaps

0;

21-MAY-1992

(first entry)

0,

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Mismatches

1 TSSEYQILTARR 12

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RESULT
R22550
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R20811
SAXAXE
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XX
                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                    Query Match
Best Local
                                                                                                                                                                                                                                                                                                                          typical features of a type I integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane-spanning domain, and a short, highly charged cytoplasmic domain. The extracellular portion contains six optential N-linked glycosylation sites and six Cys residues distributed among three C2 set Ig-related domains. A fusion protein of FCRI and a receptor ligand will be helpful to increase the potency of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      This amino acid sequence was predicted from the cDNA sequence of cDNA clone p135. It differs from the sequence predicted from clones p98/X2 and p90 (see Q21179 and Q21180, respectively) at position 25; p135 encodes a Ser residue and the other two clones predict a Thr residue. At position 58, p135 predicts Leu and p90 predicts Val. Sequences predicted from all 3 clones show the
                                              R22550;
                                                                          R22550 standard; Protein;
                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 10; Page 94a; 160pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New CD53 cell surface antigen and DNA encoding immuno-therapy and diagnosis of haematopoietic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   N-PSDB; Q21178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Seed B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GEHO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              13 'JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-JAN-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO9201049-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rapid immunoselection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human macrophage-specific FcRI receptor encoded by clone p135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      R20811;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  R20811 standard; Protein; 374 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      immunodiagnosis;
                                                                                                                                                       241 tsseyqiltarr
                                                                                                                                                                        1 TSSEYQILTARR 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1992-056864/07
                                                                                                                                                                                                                      1 Similarity
12; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Aruffo A,
                                                                                                                                                                                                                                                                                                 374 AA;
                                                                                                                                                                                                                    100.0%; ilarity 100.0%; Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              90US-0553759
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             90WO-US04986
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ction cloning technique; high affinity receptor.
                                                                                                                                                       252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Amiot M;
                                                                           374 AA
                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                    Score 57; DB 13; Pred. No. 0.0012;
                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                   DB 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cell surface antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   neoplasms, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    <u>it -</u>
                                                                                                                                                                                                                                                Length 374;
                                                                                                                                                                                                                                                                                                                                                s. A fusion protein
increase the
                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     for
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                                                                                                                                                                                                                      Gaps
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Human macrophage-specific FcRI receptor encoded by clone p90.

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                                    FIXXXXXX
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Best Local Similarity
Matches 12; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cDNA clone p90. It differs from the sequence predicted from clone p135 (see Q21178) at position 25; p135 encodes a Ser residue and clone p90 predicts a Thr residue. At position 58, p90 predicts val while clones p135 and p98/X2 (see Q21179) both predict Leu. sequences predicted from all 3 clones show the typical features of a type I integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane-spanning domain, and a short, highly charged cytoplasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and six Cys residues distributed among three C2 set Ig-related domains. A fusion protein of FCRI and a receptor ligand will be helpful to increase the potency of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Seed B, Aruffo A, Amiot M
                                                                                                                                                   Cell surface antigen; cloning; immunoselection; immunotherapy;
therapy; diagnosis; vector; FCRI; Fc receptor.
                                                                                                                                                                                                               Human FcRI (cDNA clone p135 product).
                                                                                                                                                                                                                                                      30-OCT-1996 (first entry)
                                                                                                                                                                                                                                                                                             R91438;
                                                                                                                                                                                                                                                                                                                                    R91438 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This amino acid sequence was predicted from the cDNA sequence of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 10; Page 94a; 160pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New CD53 cell surface antigen and DNA encoding it - for immuno-therapy and diagnosis of haematopoietic neoplasms,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (GEHO-) GEN HOSPITAL CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        23-JAN-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            W09201049-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rapid immunoselection cloning technique; cell surface antigen; immunodiagnosis; high affinity receptor.
Misc-difference
                                                                                                                Homo sapiens
                                                         Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                     241
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 TSSEYQILTARR 12
                                                                                                                                                                                                                                                                                                                                                                           œ
                                                                                                                                                                                                                                                                                                                                                                                                                                 tsseyqiltarr 252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Q21180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         90US-0553759
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              90WO-US04986
                                                       Location/Qualifiers 25
  58
                                    /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%;
                "amino acid 25 is Thr in clone p90 and p98 translated products"
                                                                                                                                                                                                                                                                                                                                    37'4 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 57; DB 13; Pred. No. 0.0012; Mismatches C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 374;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0,
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  DYXX
                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 12; Conservative 0
01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
18-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cDNA clone, p135 (T14717), obtd. from a cDNA library using an immunoselection cloning method. FCRI is a high affinity receptor for the Fc portion of IgG, normally located on cell surfaces of macrophages. Another isolated cDNA clone, p90 (T14719), coded for a variant (M00859) of the FCRI sequence, and a third clone, p9 (T14718), coded for an FCRI (R91439) having a different C-terminal sequence. A fusion protein of FCRI and a receptor ligand will be helpful in increasing the potency of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US5506126-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cloning of cDNA encoding cell surface antigen - useful for isolation of diagnostic and therapeutic proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      09-APR-1996
                                                                             09-APR-1996
                                                                                                                                   Homo sapiens
                                                                                                                                                           therapy; diagnosis;
                                                                                                                                                                           Cell surface antigen; cloning; immunoselection; immunotherapy;
                                                                                                                                                                                                    Human FcRI (cDNA clone p90 product).
                                                                                                                                                                                                                                 30-0CT-1996
                                                                                                                                                                                                                                                          W00859;
                                                                                                                                                                                                                                                                                  W00859 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The amino acid sequence (R91438) of human FCRI was detd. cDNA clone, p135 (T14717), obtd. from a cDNA library usir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 10; Column 55-56; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             N-PSDB; T14717.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-DEC-1992,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-FEB-1988;
                                                     25-FEB-1988;
                                                                                                          US5506126-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GEHO ) GEN HOSPITAL CORP.
                                                                                                                                                                                                                                                                                                                                                                                  1 TSSEYQILTARR 12
                                                                                                                                                                                                                                                                                                                                                        241 tsseyqiltarr 252
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Seed B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             374 AA;
                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       92US-0983647.
88US-0160416.
89US-0379076.
89US-0553759.
93US-0139273.
92US-0983647.
88US-0160416.
89US-0379076.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          88US-0160416
                                                     88US-0160416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note= "amino acid 58 is Val in p90 clone
    translated product"
                                                                                                                                                              vector; FcRI; Fc receptor
                                                                                                                                                                                                                                                                                     374
                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                                                                                          Score 57;
Pred. No.
                                                                                                                                                                                                                                                                                     Ą
                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                         0.0012;
                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 374;
                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 10
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The amino acid sequence (W00859) of human FcRI was detd. from a cDNA clone, p90 (T14719), obtd. from a cDNA library using an immunoselection cloning method. FcRI is a high affinity receptor for the Fc portion of IgG, normally located on cell surfaces of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cloning of cDNA encoding cell surface antigen of diagnostic and therapeutic proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1996-200279/20.
N-PSDB; T14719.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         for the Fc portion of IgG, normally located on cell surfaces of macrophages. Other cDNA clones (see also T14717-18) coded for variants (R91438-39) of the FcRI sequence. A fusion protein of FcRI and a receptor ligand will be helpful in increasing the potency of antibodies in therapy:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (GEHO ) GEN HOSPITAL CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13-JUL-1990;
18-OCT-1993;
                                                                                                                                                                                                                                                                                                    Fc receptor I; FcRI;
cloning.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 10; Column 55-56; 79pp; English.
                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                         Human Fc receptor I
                                                                                                                                                                                                                                                                                                                                                                  07-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                          W80448;
                                                                                                                                                                                                                                                                                                                                                                                                                 W80448 standard; Protein; 374 AA.
                                                                                                                      Misc-difference
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nilarity 100.0%;
Conservative 0
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93US-0139273.
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159
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141
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77
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103
                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                    cell surface antigen; lymphocyte; human;
                                                                                                                                   "encoded by
                                                                                                                                                          "encoded by CTG"
                                                                                                                                                                                                      "encoded by CTC"
                                   "encoded by CCC"
                                                                                    "encoded
                                                                                                                                                                                  "encoded by GAC"
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              "encoded by
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 57; DB 17;
Pred. No. 0.0012;
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                                                                                                                                    ccc.
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                                                                                     TCC'
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              GGC"
                                                              CAA"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 374;
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W97833 W97833 ID W9' XX AC W9'

W97833 standard;

Protein; 374 AA:

11

W97833;

QΥ 밁

241 tsseyqiltarr

252

1 TSSEYQILTARR 12

Query Match Best Local S Matches 12

l Similarity 100 12; Conservative

0

100.0%;

Score 57; DB 19; Pred. No. 0.0012;); Mismatches 0;

Length 374; Indels

0;

Gaps

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XX
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13-JUL-1990;
21-MAY-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Misc-difference
                                                                                    adhesion to an antibody-coated substrate. The method is useful for the isolation and cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V63442-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated genes can be expressed in a prokaryotic or eukaryotic host cells to produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V63444) which allow the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and
                                                                                                                                                                                                                                                     expression libraries is based on transient expression of an antigen in eukaryotic cells and selection of cells expressing the antigen by adhesion to an antibody-coated substrate. The method is useful for
                                                                                                                                                                                                                                                                                                                                                                                                   This is the amino acid sequence of human Fc receptor I (FCRI), as deduced from cDNA clone p135 (see V63456) isolated using a rapid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-DEC-1992;
25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US5830731-A
                                                                                                                                                                                                                                                                                                               were also obtained.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             human lymphocyte antigenic sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GEHO ) GEN HOSPITAL CORP
Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      N-PSDB;
                                                                                                                                                                                                                                                                                                           from a single patient undergoing extracorporeal interleukin-2 induction therapy. Clones p90 (see x07372) and p98 (see x07373) were also obtained. A novel method for cloning cDNAs from mamma
                                                     treatment of immune-mediated
                                                                                                                                                                                                                                                                                                                                                                                immunoselection cloning method from a cDNA library expressed
                                                                      .mmunotherapeutic applications,
                                                                                                                                                                                                                                                                                                                                                           The cDNA library was constructed from polyA RNA of cells
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                                    including
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Seed
374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Column 53-54;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 88US-0160416.
89US-0379076.
90US-0553759.
97US-0861205.
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256
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176
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                                                         infections,
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fections, diseases, and diso
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                                                               disorders of
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This is the amino acid sequence of human Fc receptor I (FcRI), as deduced from cDNA clone p90 (see X07372) isolated using a rapid immunoselection cloning method from a cDNA library expressed in COS cells. The cDNA library was constructed from polyA RNA of cells from a single patient undergoing extracorporeal interleukin-2 induction therapy. Clones p135 (see V63456) and p88 (see X07373)
                                                                                                                                                                                                                                                                                                       25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
                                                                                                                                           New cloning vector and polylinker - based on efficient cloning and expression of mammalian human lymphocyte antigenic sequences
             induction therapy.
were also obtained
                                                                                                                  Example 10; Column 53-54; 75pp; English.
                                                                                                                                                                                                WPI; 1998-609251/51.
N-PSDB; X07372.
                                                                                                                                                                                                                                       Aruffo A,
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- 85
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89US-0379076.
90US-0553759.
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77
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141
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103
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176
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expression of an antigen
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            from mammalian
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RESULT
W97834
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Best Local Similarity
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Misc-difference 141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     genes can be expressed in a prokaryotic or eukaryotic host cells to produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V63444) which allow the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            in eukaryotic cells and selection of cells expressing the antigen adhesion to an antibody-coated substrate. The method is useful for the isolation and cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V63442-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated
                                                                                                                                                             Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                          Human Fc receptor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       W97834;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            W97834 standard; Protein;
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                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                           cloning
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                                                                                                                                                                                   Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   immunotherapeutic applications, including the diagr treatment of immune-mediated infections, diseases,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 TSSEYQILTARR 12
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| 241 tsseyqiltarr 252
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171
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Pred. No. 0.0012;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    and disorders of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              be expressed and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         is useful for
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В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This is the amino acid sequence of human Fc receptor I (FCRI), as deduced from cDNA clone p98/X2 (see X07373) isolated using a rapid immunoselection cloning method from a cDNA library expressed in COS C cells. The cDNA library was constructed from polyA RNA of cells induction therapy. Clones p135 (see V63456) and p90 (see X07372) C were also obtained. A novel method for cloning cDNAs from mammalian C expression libraries is based on transient expression of an antigen by a clearly obtained. A novel method for cloning cDNAs from mammalian C expression to an antibody-coated substrate. The method is useful for the isolation and cloning of any protein which can be expressed and C transported to the cell surface membrane of a cukaryotic cell. It has been used to clone genes (see V63442-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated C genes can be expressed in a prokaryotic or cukaryotic host cells to produce the encoded protein. The invention also provides high C efficiency expression vectors (see V6341 and V6344) which allow the generation of very large mammalian expression libraries. The immunodiagnostic and immunotical and inventors is and mammalian including the diagnostic and immunotical and continued the proteins are useful for immunodiagnostic and immunotical and including the diagnostic and immunotical and including the diagnostic and immunotical and im
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 100.0%; Best Local Similarity 100.0%; Matches 12; Conservative 0;
                   Macrophage; FcRI; cell surface antigen; human; immunoselection; panning; immunodiagnosis; diagnosis; immunotherapy; gene therapy; immune disorder; infection; asthma; immune-complex disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
                                                                                                                                     Human macrophage-specific FcRI.
                                                                                                                                                                                             19-DEC-2000
                                                                                                                                                                                                                                                                                                    Y96134 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1998-609251/51.
N-PSDB; X07373.
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21-MAY-1997;
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reatment of immune-mediated in;
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                                                                                                                                                                                       (first entry)
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88US-0160416.
89US-0379076.
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97US-0861205
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Pred. No. 0.0012;
; Mismatches 0;
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parasitic
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disease;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 374;
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Fc receptor
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Francisco Residente de la companya d
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                                                                                                                             01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
23-MAR-1990;
13-JUL-1990;
                         Stamenkovic
                                                                        (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                            28-OCT-1998;
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88US-0160416. 89US-0379076. 90US-0498809. 90US-0553759.

92US-0983647 98US-0181612

WPI; 2000-586382/55 N-PSDB; A50592.

physical selection of cells expressing the antigen by adhesion to (panning on) an antibody-coated substrate such as a culture dish. CSA nucleic acids isolated by the method of the invention, and the proteins they encode, are useful for immunodiagnostic and immunotherapeutic applications, including the diagnosis and treatment of immune-mediated infections, diseases, and disorders in animals, including humans. These disorders include asthma,

diseases, and disorders in

immune-complex disease, amyloidosis, parasitic diseases or multiple sclerosis. FCRI is a high affinity receptor for the FC portion of IgG, normally located on the cell surfaces of macrophages. The ability to interfere with such bonding, or to cause it to occur on surfaces other than macrophages, is useful in therapy. A fusion protein of FCRI and a receptor ligand will be helpful to increase

The present sequence is that of a human macrophage specific as deduced from cDNA clone pl35 (see A50592), which was isol

as deduced from cDNA clone p135 (see A50592), which was isolated from a cDNA library expressed in COS cells using a novel method o

based upon transient expression of a CSA in eukaryotic cells and

the invention designed to isolate CSA nucleic acids.

The method is

of.

Example 10; Column 53-55; 75pp; English.

Isolated nucleic acid molecule encoding the CD19 cell surface antigen, useful for immunodiagnosis and immunotherapy of immune-mediated infections or disorders, e.g. asthma, immune-complex disease, parasitic

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В
RESULT 14
                                                                                                                                                                                                       PP1006183-A1
                                                                                                                                                                      07-JUN-2000
                                                                                                                                                                                                                                                                                        Human; high affinity Fc receptor; FcgammaRI; immunoglobulin; infection; immune response; CD64; monocyte; macrophage; neutrophil; eosinophil; HIV; IgG; immunosuppressive; antirheumatic; cytostatic; antiinflammatorv; AIDS; SLE; multiple myeloma; rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                               Human high affinity Fc receptor, FcgammaRI
                                                                                                                                                                                                                                                                                                                                                                                                                   11-SEP-2000 (first entry)
                                                                                                 03-DEC-1998;
                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                          systemic lupus erythematosus; tumour.
            N-PSDB; A27466
                              WPI; 2000-367968/32
                                                                                                                                    03-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Y96226 standard; Protein; 374 AA
                                                             (PLAC ) MAX PLANCK GES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             241
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          tsseyqiltarr 252
                                                                                                 98EP-0122969
                                                                                                                                  98EP-0122969
                                                               FOERDERUNG WISSENSCHAFTEN.
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RESULT 15
W86195
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local :
                                                         23-MAR-1990;
13-JUL-1990;
07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       pathways are activated, leading to immune response. The present sequence may be modified to produce recombinant versions. The recombinant recombinant for receptor consist only of the extracellular portion of the receptor and are not glycosylated i.e. they do not have transmembrane domains or signal peptides. The recombinant proteins may be used in immunoassays to determine the immune status of patients with chronic diseases of the immune system, e.g. AIDS, systemic lupus erythematosus (SLE), multiple
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   W86195 standard; Protein; 377 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present sequence is the imman first play an important regammaRI is also known as CD64. Frequences play an important role in defending the body against infections. First, pathogens are opsonised by serum immunoglobulins. The resulting complex then binds opsonised by serum immunoglobulins. The resulting complex then binds opposite the present of t
                                                                                                                                                                          01-DEC-1992;
25-FEB-1988;
                                                                                                                                                                                                                                                                      07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                       US5849898-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Fc receptor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            W86195;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  myeloma (MM), or rheumatoid arthritis. In addition, pharmaceutical compositions containing recombinant proteins may be used to treat or prevent autoimmune diseases, allergies or tumours, especially AIDS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 monocytes and macrophages, but expression can also be induced on neutrophils and eosinophils. Upon Fc receptor activation, immune effector
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence is the human high affinity Fc receptor,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel Fc receptor lacking transmembrane domains, a signal peptide, glycosylation, useful for diagnosing and treating immune disorders
                                                                                                                                                                                                                                                                                                                                15-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                rheumatoid arthritis
(GEHO ) GEN HOSPITAL CORP
                                                                                                                                               13-JUL-1989
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   l Similarity
12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                               92US-0983647.
88US-0160416.
89US-0379076.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CD36; cell surface antigen; human; cDNA library
                                                             95US-0485447
                                                                                                                       90US-0498809
                                                                                                                                                                                                                                                                      95US-0485447
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note= "this sequence contains a considerable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-28;
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                                                                                                                                                                                                                                                                                                                                                                                                                                        number of differences from the deduced from the DNA sequence the specification"
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Pred. No.
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1 TSSEYQILTARR 12

Matches 12; Query Match Best Local Similarity

Conservative

0;

100.0%;

Score 57; DB 21; Pred. No. 0.0012; Mismatches

DB 21;

Length 374; Indels.

0;

Gaps

Sequence

374 AA;

the potencies of antibodies in therapy.

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This polypeptide human Fc receptor I (FCRI). FCRI cDNAs (see V81213) were isolated using a rapid immunoselection cloning method from a cDNA library expressed in COS cells. The cDNA library was constructed from polyA RNA of cells from a single patient undergoing extroorporeal interleukin-2 induction therapy. DNA sequence analysis revealed that the cDNAs encoded type I integral membrane proteins with 3 extracellular immunoglobulin domains. The invention provides a novel method for cloning cDNAs from mammalian expression libraries. This is based on transient expression of an antigen in eukaryotic cells and physical selection of cells expressing the antigen by adhesion to an antibody-coated substrate. The method is useful for the isolation and molecular cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. CD40_cDNA (see V81198) is specifically claimed.
                                                                                                                            Query Match 100.0%; 9
Best Local Similarity 100.0%; 1
Matches 12; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cDNA encoding human CD40 antigen - useful for cloning cDNA encoding cell surface antigens, constructing cDNA libraries, expression vectors for expression in eukaryotic cells or their fragments
                                                                                                                                                                                                                                                                                          Sequence
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1 TSSEYQILTARR 12
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244 tsseyqiltarr 255
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), Seed B, Simmons D,
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; Pred. No. 0.0012;
; ""matches 0;
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D, Stamenkovic I, Stengelin S;
                                                                                                                                                                                        Length 377;
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Search completed: June Job time: 203 sec 4, 2001, 12:13:30

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Result
No.
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Maximum DB
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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      Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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57
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Gapop 10.0 , Gapext 0.5
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/cgn2_6/ptodata/2/iaa/PCTUS_COMB.pep:*
/cgn2_6/ptodata/2/iaa/backfiles1.pep:*
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          PCT US95-16311-1
US-08-469-486-52
US-08-469-688-52
US-09-036-987A-7
US-08-986-217-3
US-09-154-874-9
US-09-170-670-5
US-09-170-670-5
US-09-183-412-5
US-08-683-838A-13
US-08-683-838A-13
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US-08-375-799-15

US-08-75-299-15

US-09-090-793-9

US-09-032-372-1

US-08-240-355-409-16

US-08-372-652-2

US-08-372-652-2
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US-08-372-183-2
US-09-469-721-2
PCT-US95-17023-2
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US-08-372-652-1
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3.925 Million cell updates/sec
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Query Match
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ALIGNMENTS

JS-08-332-562A-94 Patent No. INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 91 amino acids TYPE: amino acid REFERENCE/DOCKET NUMBER: 29,768 REFERENCE/DOCKET NUMBER: 54; TELECOMMUNICATION INFORMATION: TELEPHONE: (202)672-5300 TELEFAX: (202)672-5399 TELEFAX: 000106 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS ATTORNEY/AGENT INFORMATION: CLASSIFICATION: 435 PRIOR APPLICATION DATA: SOFTWARE: Patentin Rel CURRENT APPLICATION DATA: CORRESPONDENCE ADDRESS: NUMBER OF SEQUENCES: TITLE OF INVENTION: APPLICANT: TYPE: amino aci STRANDEDNESS: s NAME: BENT, Stephen A REGISTRATION NUMBER: APPLICATION NUMBER: US 07/896,457 FILING DATE: 27-MAY-1992 FILING DATE: APPLICATION NUMBER: COUNTRY: ADDRESSEE: INFORMATION: 20007-5109 Washington 904136 Application US/08332562A E: Foley & Lardner Patentin Release #1.0, Version #1.30 HULETT, MAIK D. ENTION: FC RECEPTOR FOR IMMUNOGLOBULIN HOGARTH, BONADONNA, MCKENZIE, Ian F.C. linear 31-OCT-1994 single Bernadette M Margaret L Mark 136 US/08/332,562A 94: 54270/119/GRHA Suite 500

63.2%; Score/36;

DB 2;

Length 91;

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RESULT 3
US-08-752-929-15
; Sequence 15, Ap
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US-08-375-709-15
Sequence 15, Application US/08752929 Patent No. 5798259
                                                                                                                                                Matches
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Sequence 15, APP---
No. 568389F
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Matches 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ENERAL INFORMATION:
                                                                                                                                                                                                                                     TOPOLOGY: 11
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 2004 amino aci
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FILING DATE: 15-MAY-1992
ATTORNEY/AGENT INFORMATION:
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                                                                                     905 TASEQQALTAR 915
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                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (202)672-5300
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PPLICANT: KONDO, Kiyosi
ITLE OF INVENTION: Gene
                                                                                                                                                                                                                                                                                                                                           TELEFAX:
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REFERENCE/DOCKET NUMBER: 53
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 0 FILING DATE: 14-MAY-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                Conservative
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                                                                                                                                                                                                                                      protein
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                                                                                                                                                              63.28;
72.78;
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Synthesizing Enzymes and Process for Production of
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                                                                                                                                                             Score 36; DB Pred. No. 64;
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US-09-090-793-9
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                                                                                                                                                               Sequence 9; Application No. 6140486
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                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                APPLICANT: Calgene, LLC
TITLE OF INVENTION: Production of polyunsaturated fatty acids by expression
TITLE OF INVENTION: of polyketide-like synthesis genes in plants
FILE REFERENCE: CGNE.131.010S
CURRENT APPLICATION NUMBER: US/09/09,793
CURRENT FILING DATE: 1998-06-04
EARLIER APPLICATION NUMBER: 60/048,650
EARLIER FILING DATE: 1997-06-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/375,709
FILING DATE: 20-JAN-1995
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 53
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                  905 TASEQUALTAR 915
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ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US 0 FILING DATE: 14-MAY-1993
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FILING DATE: 15-MAY-199;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/752,929
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE:
                                                                                                                                                                                                                                                                                                                     1 TSSEYQILTAR 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20007-5109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Washington
                                                                                                                                                                                       Application US/09090793
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WEGNER, Harold C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          E: Foley's Lardner 3000 K Street, N.W.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2004 amino acids
                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           KATO, Seishi
KONDO, Kiyosi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                YAMADA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PatentIn Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-NOV-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Akiko
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kazunaga
                                                                                                                                                                                                                                                                                                                                                                          63.2%; Score 36; DB 72.7%; Pred. No. 64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Enzymes and Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gene Coding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  JP 4-147945
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US 08/178,251
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      for Eicosapentaenoic Acid Synthesizing Process for Production of Eiscosapentaenoic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                           DB 1; Length 2004;
                                                                                                                                                                                                                                                                                                                                                        Indels
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Query Match
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; ORGANISM: Shewanella putrefaciens
US-09-090-793-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 66
SOFTWARE: Patentin Ver. :
                                                                                 STRANGE IN TOPOLOGY: LINE IMMEDIATE SOURCE: LIBRARY: SYNOR/
                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Billings, Lucy J
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-855-0555
                                                                                                                                                                                                         NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ENGTH: 2004
                                                                                                                                                                                            SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: FastSEQ for CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     905 TASEQQALTAR 915
                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET:
                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 94304
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TLE OF INVENTION: CELL CYCLE RELATED PROTEINS MBER OF SEQUENCES: 13
                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 TSSEYQILTAR 11
                                                                                                                                                                               ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Palo Alto
                                                                                                                                                               amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CA
                                                                                                                                                                            395 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           E: Incyte Pharmaceuticals, Inc.
3174 Porter Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       USA
                                                                                                          SYNORAB01
                                                                                                                                                                                                                                      650-845-4166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Lal, Preeti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Corley, Neil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hillman,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Guegler, Karl J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Yue, Henry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bandman, Olga
                                                                                                                                      linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                              IBM Compatible
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                                                                                                                                                 single
                        57.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                  for Windows Version 2.0
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          Score 33; DB 3; Pred. No. 40; 2; Mismatches
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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                                    Length 395;
          Indels
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       Gaps
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Query Match
Best Local Similarity
"~+~hes 6; Conserv:
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MOLECULE TYPE: protein

US-08-240-357-2
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                                                                                                                                                                                                         Sequence 16, Application US/09335409 Patent No. 6121029
                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Cyr. Devon
APPLICANT: Goerlach, Joern
TITLE OF INVENTION: GENES FOR THE BIOSYNTHESIS OF EPOTHILONES
FILE REFERENCE: 4-30582A
CURRENT APPLICATION NUMBER: US/09/335,409
CURRENT APPLICATION NUMBER: US/09/335,409
CURRENT FILING DATE: 1999-06-17
NUMBER OF SEO ID NOS: 30
                                                                                                                     APPLICANT: Molnar, Istvan
APPLICANT: Zirkle, Ross
                                                                                                                                                       APPLICANT: Schupp, Thomas APPLICANT: Ligon, James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                               1126 TTDLYQVLAAKR 1137
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CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Gordon, Alan M. REGISTRATION NUMBER: 30 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/240,357 FILING DATE: 10-MAY-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: One Cyanamid Plaza CITY: Wayne
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                 1 TSSEYQILTARR 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 TSSEYQILTA 10
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5. 5578481
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         USA
                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               201-831-3244
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 , Alan M.
, 30,637
                                                                                                                                                                                                                                                                                                                                                                                                  57.9%;
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Cardiac Adenylyl Cyclase
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                  Length 1165;
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; TYPE: PRT ; ORGANISM: Sorangium cellulosum US-09-335-409-16

SEQ ID NO 16. LENGTH: 185

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Query Match
Best Local Similarity
Thes 7; Conserv
                              PCT-US95-16311-2
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US-08-372-652-2
                                               RESULT
Sequence 2, Application PC/TUS9516311 GENERAL INFORMATION:
                                                                                                                                                                  Matches
                                                                                                                                                                                Query Match
Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                              NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOSTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/372/652
FILLING DATE: 13-JAN-1995
                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 451 amino acids
                                                                                               357 TQEEYALLTA 366
                                                                                                                                                                                                                                                                MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 13-JAN-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
TELEFAX: 617/542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORRESPONDENCE ADDRESS:

ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street, Suit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Seol, Wongi
APPLICANT: Choi, Hueng-Sik
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ITLE OF INVENTION: RETINOID X RECEPTOR-INTERACTING METHODS AND METHODS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       158 ASDQQILTALR 168
                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: USA
ZIP: 02110-2804
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STATE: MA
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TELEX: 20015
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                                                                                                                                TSSEYQILTA 10
                                                                                                                                                               l Similarity
6; Conserv
                                                                                                                                                                                                                                                                                                                amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                 linear
                                                                                                                                                                                                                                                              protein
                                                                                                                                                                                                                                                                                             not relevant
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                                                                                                                                                                                 Score 31; DB 2;
Pred. No. 1.2e+02
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Pred. No. 43;
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                                                                                                                                                                Mismatches
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MOLECULE TYPE:

PCT-US95-16311-2
                                                                                                                                                                                                                                                                               Sequence 2, Application Patent No. 6005086 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR DATE:
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: 08/372,652
FILING DATE: 13-JAN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Clark, Paul T:
NAME: Clark, Paul T: 30,162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                 TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: bi,
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
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                                                                                                                                                                                                                                                                                                                                                                                                                                    1 TSSEYQILTA 10
                                                                                                                                                                                                                                                                                                                                                                                                     357 TQEEYALLTA 366
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY
                                                                                                   STREET: 444 South
CITY: Los Angeles
STATE: CA
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COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER:
                                                                                                                              ADDRESSEE: Pretty, Schroeder, Brueggemann & Clark STREET: 444 South Flower Street, Suite 2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Boston
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                                                                                 COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 451 amino acids
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                                                                   90071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    amino acid
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                                                                                   USA
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                                                                                                                                                                                                Evans, Ronald M.
Forman, Barry M.
Weinberger, Cary A.
WEINDERGER, CARY A.
VENTION: METHOD FOR MODULATING PROCESSES MEDIATED
VENTION: BY FARNESOID ACTIVATED RECEPTORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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compatible
PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   54.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08/372,652
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Pred. No. 1.2e+02;
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Query Match
Best Local Similarity
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                                                             TELEFAX: 619-546-9392
NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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SEQUENCE CHARACTERISTICS:
LENGTH: 469 amino acid
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                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
                                                                                                                                   TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                   CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Weinberger, FITLE OF INVENTION: MET
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 OLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               UMBER OF SEQUENCES:
                                                                                                                                             NAME: Reiter, Stephen E.
REGISTRATION NUMBER: 31,192
REFERENCE/DOCKET NUMBER: P41 9844
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CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 469 amino acids TYPE: amino acid TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Reiter, Stephen E.
REGISTRATION NUMBER: 31,192
REFERENCE/DOCKET NUMBER: P4
                                                                                                                 TELEPHONE:
                                                                                                                                                                                                                                                                                          FILING DATE
                                                                                                                                                                                                                                                                                                                                       SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                     APPLICATION NUMBER:
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                                               ENGTH:
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                             amino acid
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619-546-9392
                                               469 amino acids
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                                                                                                                                                                                                                                                                                                                                     PatentIn Release #1.0, Version #1.25
                                                                                                                 619-546-4737
                                                                                                                                                                                                                                                                                                                                                                                     Floppy disk
protein
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Pred. No.
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US-09-469-721-2
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Best Local Similarity
Matches 6; Conserv
                                                                                           Sequence 2, Applic Patent No. 5728548
                                                                                                                                                                                                                                                 Matches
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Best Local Similarity
                                                                           GENERAL INFORMATION:
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                                                                                                                                                                                     375 TQEEYALLTA 384
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LENGTH: 469 amino acids
TYPE: amino acid
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                                           TITLE OF INVENTION: STEROI
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NAME: Relier, Stephen E.
REGISTRATION NUMBER: 31,192
REFERENCE/DOCKET NUMBER: P4:
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                              NUMBER OF SEQUENCES:
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CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM:
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                                                                                                          Application US/08496631
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VENTION: METHOD FOR MODULATING PROCESSES MEDIATED
VENTION: BY FARNESOID ACTIVATED RECEPTORS
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Genetics Institute,
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                                                                                                                                                                                                                                                               54.48;
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                                             STEROID RECEPTOR RR1
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Pred. No.
                                                                                                                                                                                                                                                                 Score 31;
Pred. No. 1
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STREET: o, cambridge

87 CambridgePark Drive

Massachusetts

COUNTRY:

U.S.A.

02140

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Query Match
Best Local Similarity
Ches 6; Conserve
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US-08-372-652-1
                     COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/372,652

FILING DATE: 13-TAN-1995

ATTORNEY/AGENT INFORMATION:

REGISTRATION NUMBER: 30,162

REFERENCE/DOCKET NUMBER: 00786/246001

TELECOMMUNICATION INFORMATION:

TELEPHONE: 617/542-5070
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  tent No. 5932699
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Brown, Scott A.
REGISTRATION NUMBER: 32,724
REFERENCE/DOCKET NUMBER: GIT
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 498-8224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                      UMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      378 TQEEYALLTA 387
                                                                                                                                                                                                                                                                               STATE: MA
COUNTRY: USA
ZIP: 02110-2804
                  TELEFAX:
                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Fish & Richardson P.C. STREET: 225 Franklin Street, Suite 3100
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                  617/542-8906
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Choi, Hueng-Sik
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Pred. No. 1
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Best Local Similarity
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 484 amin -
Type
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Best Local 9
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                                                                       MOLECULE TYPE:
                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION: TELEPHONE: 617/542-5070
                                                                                                                                                                                                                                                                      FILING DATE: 13-JAN-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 02110-2804.
COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
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                                                                                                                                                                       TELEPHONE: 617/542-8906
                                                                                                                                                                                                                           NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 00
                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 08/3 FILING DATE: 13-JAN-1995
                                                                                      TOPOLOGY:
                                                                                                STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                  FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
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225 Franklin Street, Suite 3100
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Choi, Hueng-Sik
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Score 31; DB 5;
Pred. No. 1.3e+02;
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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ALIGNMENTS

RESULT A41357

Fc gamma (IGG) receptor I (high affinity) N;Alternate names: CD64

form

b - human

C;Species: Homo sapiens (man)
C;Date: 03-Apr-1992 #seguence_revision 03-Apr-1992 #text_change 21-Jan-2000
C;Accession: A41357; S03019

R;Allen, J.M.; Seed, B. Science 243, 378-381, 1989
A;Title: Isolation and expression of functional high-affinity Fc receptor complementa A;Reference number: A41357; MUID:89100284

A; Accession: A41357 A; Molecule type: mRNA A; Residues: 1-344 <ALL1>

A;Cross-references: GB:X14355; GB:M21090; NID:g31333; PIDN:CAA32536.1; PID:g31334

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RESULT . 2
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A39878
Comman (IgG) receptor I-A (high affinity) precursor - human recursor and recursor in human high affinity receptor for large recursor in human recursor in high affinity receptor for large recursor in human recursor in high affinity receptor for large recursor in high recursor in high areas in high areas in high all receptor for large recursor in high receptor recursor in high receptor for large recursor in high recursor in high areas in high areas in high receptor for large recursor in high receptor for large recursor in high recursor in high receptor for large recursor in high recursor in high recursor for large recursor in high recursor in high recursor for large recursor for 
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A; Residues: 1-344 < ALL2>
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Nucleic Acids Res.
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Res. 16, 11824, 1988
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A;Molecule type: mRNA
A;Residues: 1-336 <RE
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A;Residues: 1-24,'S',26-337,'T',339-374 <AL2>
A;Cross-references: EMBL:X14356; NID:g31331; PIDN:CAA32537.1; PID:g31332
A;Perez, C.; Wietzerbin, J.; Benech, P.D.
Mol. Cell. Biol. 13, 2182-2192, 1993
A;Title: Two cis-DNA elements involved in myeloid-cell-specific expression and gamma int
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 R;Allen,
       ;Cross-references:
                                                                                                                                                                                   c gamma (IgG) receptor high affinity mouse; Species: Mus musculus (house mouse); Date: 02-Jul-1996 #sequence_revision 02-Jul
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              \; Map position:
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                                                                                           Title: Linkage on chromosome 3 of autoi; Reference number: I48471; MUID:93242399
                                                                                                                                                      ;Prins,
                                                           Status: preliminary; translated
                                                                                Accession:
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Best Local
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Accession: I57525
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Title: Isolation and expression of functional high-affinity Reference number: A41357; MUID:89100284 Accession: B41357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cross-references: GDB:135911;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Molecule type: mRNA residues: 1-24,'S',26-337,'T',339-374 <AL1> Cross-references: GB:X14356; GB:M21091; NID:g31331;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cross-references: GB:L03418; NID:g184840; Allen, J.M.; Seed, B. ience 243, 378-381, 1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cross-references: GB:S57204; NID:g298692;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Accession: S03018
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cleic Acids Res. 16, 11824, 1988
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3-313/Domain: transmembrane #status p: 78,152,159,163,195,240/Binding site:
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                                                                                                                                                    J.B.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Invest.
                                                                                                                                                                                                                                                                                                                                                       TSSEYQILTARR
                                                                                                                                                                                                                                                                                                                                                                                           Similarity
12; Conser
                                                                                I48471
                                                                                                                               Todd, J.A.;
95-698, 1993
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larity 100.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              90,
       EMBL: X70980;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GB:M63830;
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                                                                                                                                                                                                                                                                                                                    252
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                                                                                                                                                Rodrigues, N.R.; Ghosh, S.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GB:M63835; NID:g180277; PIDN:AAA35678.1; PID:g180279 Doebele, R.; Pan, L.C.; Salmon, J.E.; Kimberly, R.P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OMIM: 146760
   NID: g311748;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nomology <IMM2>
                                                           from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                     Score 57; -DB 1;
Pred. No. 0.00063;
Mismatches 0;
                                                                                                              autoimmune
                                                                                                                                                                                 02-Jul-1996  #text_change
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  predicted <TMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 carbohydrate
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PIDN:CAA50311.1; PID:g311749
                                                                                                              diabetes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      <EXT>
                                                                                                                                                                                                                                                                                                                                                                                                                       Length 374
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                                                                                                                                              Hogarth, P.M.; Wicker, L.S.; Gaf:
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                                                                                                              and
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                                                                                                              defective
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                                    A;Cross-references: GB:AE004638; GB:AE004091; NID:g9948116; PIDN:AAG05501.1; GSPDB:GNA;Experimental source: strain PAO1
                                                                        A; Molecule type:
A; Residues: 1-40
                                                                                                        A; Status: preliminary
                                                                                                                                             A;Title: Complete genome sequence of Pseudomonas aeruginosa A;Reference number: A82950; MUID:20437337
                                                                                                                                                                                                                     adman,
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                                                                                                                             A;Accession: H83382
                                                                                                                                                                               Nature 406, 959-964,
                                                                                                                                                                                                                                    R;Stover, C.K.; Pham, X.Q.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C; Keywords: immuno F; 127-179/Domain:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C; Species: Mus musculus (house mouse)
C; Date: 18-Jun-1993 #sequence_revision 18-Nov-1994
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A; Residues: 1-404 <SEA>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Note: sequence extracted from R; Sears, D.W.; Osman, N.; Tate, J. Immunol. 144, 371-378, 1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C; Accession:
                                                                                                                                                                                                                                                                                                                                                 RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Cross-references: GB:M31314; NID:g200752; PIDN:AAA40056.1; Superfamily: Fc gamma receptor I; immunoglobulin homology.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Status: preliminary
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Molecule type: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R;Osman,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            N; Alternate
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Best Local S
Matches 8
                                                                                                                                                                                                                                                       Accession:
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ol. 148, 15
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8; Conser
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                                                                                                                                                                                                                 Brody,
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                                                                                                                                                                                                                 Erwin,
L.L.; (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 homology
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Pred. No.
                                                                                                                                                                                                                   Coulter,
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Pred.
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r, S.N.; Folger,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     <IMM>
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12;
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#text_change
                                                                                                                   Length 336;
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21-Jan-2000
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                                                                                                   Gaps
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high affinity Fc receptor high affinity NCBIN: 85208, PID:g200753 Fc gamma NCBIN: 85212, RI

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Length 404
0,
Gaps
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probable porin PA2113 [imported] - Pseudomonas aeruginosa (strain PAO1) Species: Pseudomonas aeruginosa ;Date: 15-Sep-2000 #sequence_revision 15-Sep-2000 #text_change 31-Dec-2000 ; Warrener, K.R.; Kas, PA01, an opportunistic Hickey, Larbig, M.J.; K:; L

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A;Introns: 21/1; 98/3; 130/3; 143/1; 178/3
C;Superfamily: Caenorhabditis elegans hypothetical protein C31H1.
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A;Title: Expression of the eicosapentaenoic acid
A;Reference number: 220764; MUID:97419510
A;Accession: T30185
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                                                                                                                                                                                                                                                                                                                                                                             submitted to the EMBL Data Library, December 1995
A;Description: The sequence of C. elegans cosmid C31H1
A;Reference number: Z18395
                                  S46668
                                                 RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           hypothetical protein C31H1.2 - Ca
C;Species: Caenorhabditis elegans
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nypothetical
                                                                                                                                                                                                                                                                                                                  Status: preliminary; translated from GB/EMBL/DDB: Molecule type: DNA Residues: 1-197 <LET>
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Date: 20-Sep-1999 #sequence_revision 20-Sep-1999 #text_change 04-Mar-2000
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Cross-references: EMBL:U73935; NID:g2529413; PID:g2529420; PIDN:AAB81125.1
Experimental source: strain SCRC-2738
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Species: Shewanell Date: 02-Sep-2000
                                                                                                                                                                         Query Match
Best Local
                                                                                                                                                                                                                                                                                                  Cross references: EMBL: U42848; NID: g1125817; PID: g1125818; PIDN: AAA83606.1; CESP: C31H
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Best Local Similarity
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Best Local
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                                                                                               TSSEIQLLSMKR 34
                                                                                                                           TSSEYQILTARR 12
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                yeast (Saccharomyces cerevisiae)
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50.0%;
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58.3%;
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    Caenorhabditis elegans

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Pred. No.
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R; Margot, P.: Mauel, C.: Karamata, D.

Rol. Microbiol. 12, 535-545, 1994
A; Title: The gene of the N-acetylglucosaminidase,
A; Reference number: S49389; MUID:95020588
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Species: Bacillus subtilis
C;Date: 15-Feb-1996 #sequence_revision 01-Mar-1996
C;Accession: S60137; S49390; F69654
                                                                                                                                                                   R;Kunst, F.; Ogasawara, N.; Moszer, I.; Albertini, A.M.; Alloni, G.; Azevedo, V.; Be
C.; Bron, S.; Brouillet, S.; Bruschl, C.V.; Caldwell, B.; Capuano, V.; Carter, N.M.;
A.; Ehrlich, S.D.; Emmerson, P.T.; Entian, K.D.; Errington, J.; Fabret, C.; Ferrari,
Nature 390, 249-256, 1997
                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Residues: 1-880 <MAR>
A; Cross-references: EMBL: U02562; NID: g476091;
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C;Genetics:
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Mol. Cell. Biol.
                                                                                                                                                 A; Authors: Foulger, D.; Fritz, C.; Fujita, M.; Fujita, Y.; Fuma,
                                                                                                                                                                                                                                                                                                                               A; Accession: S49390
                                                                                                                                                                                                                                                                                                                                                                                                                              A; Experimental source: strain 168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Reference number:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             submitted to the EMBL Data Library,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Description:
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Residues: 1-433 <LET>
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Residues: 1-433 <HUB>
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A;Cross-references: GB:AL009198; GB:AL123456; NID:g3242262; PIDN:CAA15760.1;
A;Experimental source: strain H37Rv
                                                                                                  A;Authors: Sqares, R.; Súlston, J.E.; Taylor, K.; Whitehead, S.; Barrell A;Title: Deciphering the biology of Mycobacterium tuberculosis from the A;Reference number: A70500; MUID:98295987
A;Accession: F70972
                                                                                                                                                                                                                                                       C;Species: Mycobacterium tuberculosis
C;Date: 17-Jul-1998 #sequence_revision 17-Jul-1998 #text_change 20-Jun-2000
C;Accession: F70972
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                                             A; Residues:
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F;1-27/Domain: signal sequence #status predicted
F;28-880/Product: beta-N-acetylhexosaminidase #st
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A;Experimental source: cultivar Co
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                                                               ;Molecule type: DNA
                                                                                                                                                                              ajandream, M.A.; Rogers,
ature 393, 537-544, 1998
                                                                                                                                                                                                                  Connor, R.; Davies, R.; Parkhill, J.; Garnier, T.; Churcher, C.; Harris, D.; Connor, R.; Davies, R.; Devlin, K.; Feltwell, T.; Gentles, S.; Hamlin, N.; Ho
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                                                                             Status: preliminary; nucleic acid sequence not shown; translation not shown
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Residues: 1-277 <BEV>
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257 TSTNYKILNASR 268
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                                                                                                                                                                                                                                                                                                                   Mycobacterium tuberculosis (strain H37RV)
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                                                                                                                                                                                                    J.; Rutter,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           52; GSPDB:GN00062; ATSP:F27B13.170 Columbia; BAC clone F27B13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 34;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score
Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; Wambutt,
May 1999
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                                                                                                                                                                                                  s.;
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                                                                                                                                                         Barrell, B.G.
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                  PID: 9266165
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123 SSVYQIRRARR 133
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C; Genetics: A; Gene: PA3
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A; Residues: 1-238 <STO>
A; Cross-references: GB:AE004747; GB:AE004091; NID:g9949362;
                                                                                                                                                                                                                                                                                                       Nature 406, 959-964, 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Gene: amiD
C; Superfamily:
                                                                                                                                                             A; Experimental source: strain PAO1
                                                                                                                                                                                                                                                   A;Reference number: A82950;
A;Accession: C83240
                                                                                                                                                                                                                                                                                                                                          adman, S.; Yuan,
                                                                                                                                                                                                                                                                                                                                                         R;Stover, C.K.; Pham, X.Q.; Erwin,
                                                                                                                                                                                                                                                                                                                                                                                                           probable transcription regulator PA3249 [imported]
C; Species: Pseudomonas aeruginosa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Map position:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Gene: NCSP:B23L21.220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C; Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Molecule type: DNA
A; Residues: 1-863 <SCH>
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A; Accession: T49709
                                                                                                                                                                                                                                                                                       A; Title: Complete genome
                                                                                                                                                                                                                                                                                                                                                                            C; Accession:
                                                                                                                                                                                                                                                                                                                                                                                         C;Date: 15-Sep-2000 #sequence_revision 15-Sep-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Experimental source:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Cross-references: EMBL:AL356172;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          submitted to the Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          R; Schulte, U.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C; Accession: T49709
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C;Date: 02-Jun-2000 #sequence_revision 02-Jun-2000 #text_change 09-Jun-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C; Species: Neurospora crassa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         related to glucan 1,3-beta-glucosidase precursor [imported] - Neurospora crassa
N;Alternate names: protein B23L21.220
                                                                                                                                                                                                                                 A;Status: preliminary
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Best Local S
Matches 6
                                                 Query Match
Best Local Similarity
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Best Local Similarity 77.0
Matches 7; Conservative
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2 SSEYQILTARR 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TSSEYQILTARR
                                                                                                                                                                                                                                                                                                                  uan, Y.; Brody, L.L.; Coulter, S.N.; Folger, K.R.; Olson, M.V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
6; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Aign, V.; Hoheisel, J.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                    Conservative
                                                                                                                                                                                                                                                                    sequence of Pseudomonas aeruginosa PAO1, an opportunistic 50; MUID:20437337
                                                     57.9%;
72.7%;
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77.8%;
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                                   Score 33; DB Pred. No. 35; 0; Mismatches
                                    0,
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B23L21; strain OR74A
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Pred. No.
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Pred. No. 45;
0; Mismatches
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    Pseudomonas aeruginosa (strain

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                                                                    Length 238;
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                                   Gaps
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K.; L
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C;Species: Streptomyces coelicolor
C;Date: 03-Dec-1999 #sequence_revision 03-Dec-1999 #text_change 31-Jan-2000
C:Date: 05-Dec-1999 #sequence_revision 03-Dec-1999 #text_change 31-Jan-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Gene: trpC2; SCOEDB:SCE8.04c
C;Superfamily: indole-3-glycerol-phosphate synthase; trpC homology
                                                                                                                                                                                                                A; Note: DKFZp762M186.1
                                                                                          Query Match 57.9
Best Local Similarity 70.0
Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                      Bloecker, H.; Boecher, M.; Brandt, P.; Mewes, H.W.; Weil, B.; Wiemann, S.; Bhmitted to the Protein Sequence Database, June 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         pothetical protein DKFZp762M186.1 - human (fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 57.9%;
Best Local Similarity 87.5%;
Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Saunders, D.; Harris, D.; Bentley, S.D.; Parkhill, J.; Barrell, B.G.; Rajandream, M.A. bmitted to the EMBL Data Library, March 1999 Reference number: Z21604 Accession: T36303
                                                                                                                                                                                                                                                                                                      Status: preliminary
Molecule type: mRNA
Residues: 1-271 <AAA>
                                                                                                                                                                                                                                                                                                                                                                           Reference number: Z25143
Accession: T50620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Species: Homo sapiens (man)
Date: 21-Jul-2000 #sequence_revision 21-Jul-2000 #text_change 21-Jul-2000
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                                                                                                                                                                                                                                          Genetics:
                                                                                                                                                                                                                                                    Cross-references: EMBL:AL359590
Experimental source: adult melanoma (MeWo cell line); clone DKFZp762M186
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cross-references: EMBL:AL035654; PIDN:CAB38582.1; GSPDB:GN00070; SCOEDB:SCE8.04c Experimental source: strain A3(2)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Accession: T50620
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117 TRAEYQILLA 126
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                                                                                                                  57.9%;
70.0%;
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                                                                                             1; Mismatches
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Pred. No. 40;
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Perfect score:
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1 TSSEYQILTARR 12
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Gapop 10.0 , Gapext 0.5
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Copyright (c) 1993 - 2000 Compugen Ltd.
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 NIFK_KLEPN
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P26151 mus musculu
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Best Local Similarity 100 Matches 12; Conservative
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P26151;
01-MAY-1992 (Rel. :
01-MAY-1992 (Rel. :
15-JUL-1999 (Rel. :
HIGH AFFINITY IMMUN
                                                     "Structure and mapping of the gene encoding mouse gamma RI and chromosomal location of the human Fc J. Inmunol 148:1570-1575(1992).

1 TOWCTION: BINDS TO THE FC REGION OF IMMUNOGLOB ARFTINTY PROFESSION.
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Sears D.W., Osman N., 7
"Molecular cloning and
receptor for IgG.";
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Mus musculus (Mouse).

Chordata;
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SUBCELLULAR LOCATION: TYPE I MEMBRANE PROTEIN
TISSUE SPECIFICITY: MACROPHAGE-SPECICIC.
SIMILARITY: CONTAINS 3 IMMUNOGLOBULIN-LIKE C2.
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IMMUNOGLOBULIN GAMMA FC RECEPTOR I PRECURSOR (FC-GAMMA PRICEPTOR I).
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                                      / AB972;
Andrews S.,
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EMBL; M31314; AAA40056.1;
PIR; A43511; A43511.
PIR; A46480; A46480.
                                                                                                                                                                                                                                                                                                                                                                              use by non-profit institutions as long modified and this statement is not removed entities requires a license agreement (See
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IG-LIKE C2-TYPE DOMAIN
IG-LIKE C2-TYPE DOMAIN
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Saccharomyces cerevisiae (Baker's yeast).
Eukaryota; Fungi; Ascomycota; Saccharomyc
                         MEDLINE=94158870; PubMed=8114728;
Hubbard E.J.A., Jiang R., Carlson
"Dosage-dependent modulation of g
                                                                                    SEQUENCE FROM N.A.
                                                                                                                 Saccharomycetales;
NCBI_TaxID=4932;
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Rashid M.H., Mori M., Sekiguchi J.;
"Glucosaminidase of Bacillus subtilis: cloning, regulation, primary
structure and biochemical characterization.";
structure and biochemical characterization.";
Microbiology 141:2391-2404 (1995).
MICROBION: CELL MALL HAYDROLASE NOT INVOLVED IN CELL AUTOLYSIS. IT
HYDROLYZES THE BETA-1,4 GLYCAN BOND BETWEEN THE N-
ACETYLGLUCOSAMINYL AND THE N-ACETYLMURAMOYL RESIDUES IN-THE GLYCAN
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"Deciphering the biology of Mycobacterium tuberculosis from complete genome sequence.";
Nature 393:537-544(1998).
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                                                                                                     the European Bioinformatics Institute. The use by non-profit institutions as long modified and this statement is not removed.
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PROSITE; PS00571; AMIDASES; FALSE_NEG

Amidase;

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InterPro;

IPR000120;

PF01425;

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2_ENTHI
GIL2_ENTHI STANDARD; PRT; 288 AA.
Q03077;
Q1-JUL-1993 (Rel. 26, Created)
O1-JUL-1993 (Rel. 26, Last sequence update)
O1-NOV-1995 (Rel. 32, Last annotation update)
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Q9Z4X0;
30-MAY-2000
                                                                                                                                                                                                                                                                                                              modified and this statement is not removentities requires a license agreement (S or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                               Tryptophan biosynthesis; Lyase; Decarboxylase; Antibiotic biosynthesis.
                                                                                                                                                                                                                                                                               EMBL; AL035654; CAB38582.1; HSSP; P00909; 1PII.
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                                                                                                                                                                                                                                                           Pfam; PF00218;
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30-MAY-2000 (Rel. 39, Last annotation update)
INDOLE-3-GLYCEROL PHOSPHATE SYNTHASE 2 (EC 4.1.1.48) (IGPS).
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SIMILARITY: BELONGS TO THE TRPC FAMILY.
                                                                                                         EYQILEAR 130
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EMBL; L06065; AAA16054.1;
                                                                                                                                                                        or send an email to license@isb-sib.ch)
                                                                                                                                                                                                  use by non-profit institutions as long modified and this statement is not removed.
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MEDLINE=94043255; PubMed=8226970;
MCDCJ J.J., Mann B.J., Vedvick T.S.,
Petri W.A. Jr.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "Molecular cloning of cDNA and genomic sequences coding for the kilodalton subunit of the galactose-inhibitable lectin of pathog Entamoeba histolytica.";
                                                                                       SIGNAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tannich E., Ebert F.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Entamoebidae;
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                                                                                                                                                                                                                                                                                                                                     istolytica galactose-specific adherence lectin.",
    Biol. Chem. 268:24223-24231(1993).
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                                                                                                                                                                                                                                                                                                                                                             Structural analysis of the light subunit of the Entamoeba
                                                                                                                                                                                                                                                                                                                    Biol. Chem. 268:24223-24231(1993).
I- FUNCTION: LECTIN THAT BINDS GALACTOSE
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         GALACTOSE-INHIBITABLE LECTIN 35 KDA SUBUNIT.
S -> A (IN REF. 2).
E -> K (IN REF. 2).
E -> G (IN REF. 2).
DECF366D3DAFA24B
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RESULT
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SEQUENCE FROM N.A. STRAIN=NA1000;
                                                                                                                                 PARB_CAUCR 005190;
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                                 NCBI_TaxID=76;
                                             Caulobacter
                                                      Bacteria; Proteobacteria;
                                                                  Caulobacter crescentus.
                                                                                        CHROMOSOME PARTITIONING
                                                                                                 01-OCT-2000
01-OCT-2000
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                                                                             PARB
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(Rel. 40, Last ann
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54.5%;
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                                                    alpha subdivision; Caulobacter group;
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Pred. No.
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Query Match Best Local

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Length 288

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MEDLINE=97207014; PubMed=9054507; Mohl D.A., Gober J.W.;

cycle-dependent polar localization of chromosome partitioning

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P53581;
                                                                                               This SWISS-PROT entry is copyright: It is produced through a between the Swiss Institute of Bioinformatics and the EMBIT the European Bioinformatics Institute. There are no restrict use by non-profit institutions as long as its content is modified and this statement is not removed. Usage by and f
                                                                                                                                                                                                                                                                                                                                                                                            "Sequence analysis of the genome of the unicellular cyanobacterium Synechocystis sp. strain PCC6803. I. Sequence features in the 1 Mb region from map positions 64% to 92% of the genome.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-OCT-1996 (Rel. 34, Created)
01-OCT-1996 (Rel. 34, Last sequence update)
01-OCT-1996 (Rel. 34, Last annotation update)
PUTATIVE METHIONINE AMINOPEPTIDASE C (EC 3.4.11.18) (MAP) (PEPTIDASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         use by non-profit institutions as long modified and this statement is not removed. entities requires a license agreement (See or send an email to license@isb-sib.ch).
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EMBL; D64003; BAA10478.1; HSSP; P07906; IMAT.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sugiura M., Tabata S.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synechocystis sp. (strain PCC 6803). Bacteria; Cyanobacteria; Chroococcales;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE-96127529; PubMed-8590279;
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                                                                                                                                                                                                                                                                                                          FUNCTION: REMOVES THE AMINO-TERMINAL MI PROTEINS (BY SIMILARITY).

CATALYTIC ACTIVITY: L-METHIONYLPEPTIDE
                                                                                                                                                                                                                                               COFACTOR: COBALT (BY SIMILARITY).
SIMILARITY: BELONGS TO PEPTIDASE FAMILY M24A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FUNCTION: INVOLVED IN CHROMOSOME PARTITION. LOCALIZE TO BOTH POLES OF THE PREDIVISIONAL CELL FOLLOWING COMPLETION OF DNA REPLICATION.
                                                                                                                                                                                                                                MAP FAMILY 1.
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6; Conserv
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293 AA; 31910 MW; E7D557D90F61469C CRC64;
                                                            email to license@isb-sib.ch).
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BELONGS TO THE PARB PROTEIN FAMILY.
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                                                                                  http://www.isb-sib.ch/announce/
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RESULT
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Best Local
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                                                                                                                                                                                                                                                                                                                                                        Jones P.G., Lombardi S.J., Cockett M.I.;

"Cloning and tissue distribution of the human G protein beta 5 cDNA
Biochim. Biophys. Acta 1402:288-291(1998).

-I- FUNCTION: GUANINE NUCLEOTIDE-BINDING PROTEINS (G PROTEINS) ARE
INVOLVED AS A MODULATOR OR TRANSDUCER IN VARIOUS TRANSMEMBRANE
SIGNALING SYSTEMS. THE BETA AND GAMMA CHAINS ARE REQUIRED FOR T
GTPASE ACTIVITY, FOR REPLACEMENT OF GDP BY GTP, AND FOR G PROTE
PROSITE; PS00678; WD_REPEATS_1; 3.
PROSITE; PS0082; WD_REPEATS_2; 6.
PROSITE; PS50294; WD_REPEATS_REGION; 1.
Transducer; Repeat; WD repeat; Multigene
REPEAT 61 91 WD 1.
                                                                                                                                                                                            use by non-profit institutions as long as its cont modified and this statement is not removed. Usage by entities requires a license agreement (See http://www.or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                  This SWISS-PROT entry is copyright. It is produced throubetween the Swiss Institute of Bioinformatics and the the European Bioinformatics Institute. There are no reuse by non-profit institutions as long as its contents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CHAIN 5).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-JUL-1998 (Rel. 36, Last sequence update)
01-OCT-2000 (Rel. 40, Last annotation update)
GUANINE NUCLEOTIDE-BINDING PROTEIN BETA SUBUNIT 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-JUL-1998
01-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      014775;
15-JUL-1998
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METAL 147 COB
METAL 158 COB
METAL 221 221 COB
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                                                                                                                                                    EMBL; AF0176
MIM; 604447;
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InterPro; IPR002467;
                                                                                               InterPro; IPR001632;
InterPro; IPR001680;
Pfam; PF00400; WD40;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
                                                                     PRINTS; PR00319; GPROTEINB.
PRINTS; PR00320; GPROTEINBRPT
                                                                                                                                         HSSP; P04901; 1GP2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=9606;
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                                                                                                                                                                                                                                                                                                     EFFECTOR INTERACTION.
SUBUNIT: G PROTEINS ARE COMPOSED OF 3 UNITS (ALPHA, SUBUNIT: G PROTEINS ARE COMPOSED IN MULTIPLE TISSUES.
TISSUE SPECIFICITY: EXPRESSED IN MULTIPLE TISSUES.
SIMILARITY: CONTAINS 7 WD REPEATS (TRP-ASP DOMAINS).
                                                                                                                                                                    AF017656; AAC63826.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity 50. 6; Conservative
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo
                                                                                                                                                                                                                                                 restrictions
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                                                                                                                                                                                                                                                                                                                                 BETA
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RESULT
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Best Local Similarity
Thes 7; Conserv
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                                                                       This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way
                                                                                                                                                                                                                                                                 Betty M., Harnish S.W., Rhodes K.J., Cocke "Distribution of heterotrimeric G-protein the rat brain.";
                                                use by non-profit institutions as long a
modified and this statement is not removed.
entities requires a license agreement (See )
                                  or send an email to license agreement (s
                                                                                                                                         -!- SUBUNIT: G PROTEINS ARE COMPOSED OF 3 UNITS (ALPHA, E-!- TISSUE SPECIFICITY: BRAIN AND AT MUCH REDUCED LEVELS -!- SIMILARITY: CONTAINS 7 WD REPEATS (TRP-ASP DOMAINS).
                                                                                                                                                                                   Neutoscience 85:475-486(1998).

-i- FUNCTION: GUANINE NUCLEOTIDE-BINDING PROTEINS (G PROTEINS) ARE INVOLVED AS A MODULATOR OR TRANSDUCER IN VARIOUS TRANSMEMBRANE SIGNALING SYSTEMS. THE BETA AND GAMMA CHAINS ARE REQUIRED FOR T GTPASE ACTIVITY, FOR REPLACEMENT OF GDP BY GTP, AND FOR G PROTE FEFECTOR INTERACTION.
                                                                                                                                                                                                                                                                                                                                                                               SPECIES=Rat; STRAIN=SPRAGUE-DAWLEY; TISSUE=Brain Levay K., Slepak V.Z.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Watson A.J., Katz A., Simon M.I.;
"A fifth member of the mammalian G-protein beta-subunit family.

Expression in brain and activation of the beta 2 isotype of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GBB5_MOUSE STANDARD; PRT; 353 AA. P54314; 035354; 01-0CT-1996 (Rel. 34, Created) 01-0CT-1996 (Rel. 34, Last sequence update) 30-MAY-2000 (Rel. 39, Last annotation update) GUANINE NUCLEOTIDE-BINDING PROTEIN BETA SUBUNIT 5 (TRANSDUCIN BETA
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SEQUENCE
                                                                                                                                                                                                                                                                                                                SPECIES-Rat; STRAIN-SPRAGUE-DAWLEY; MEDLINE-98283588; PubMed-9622245;
                                                                                                                                                                                                                                                                                                                                                                     Submitted (MAY-1997) to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mus musculus (Mouse), and Rattus norvegicus (Rat). Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eukaryota; Eutheria; Rodentia; Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                           SEQUENCE OF 11-198 FROM N.A.
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70.0%;
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                                                                                                                                                                                                                                                                                  Cockett M. otein beta
                                             (See http://www.isb-sib.ch/announce/
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EMBL;

EMBL;

L34290; AZ AF001953;

AAA93084.1;

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1 TSSEYQILTA

Matches Query Match Best Local:

Local Similarity nes 7; Conserv

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Score 33; DB Pred. No. 24;

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045435;
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15-JUL-1999 (Rel. 3
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HYPOTHETICAL 49.0 K
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                                                                                  use by non-profit institutions as long modified and this statement is not removed, entities requires a license agreement (See
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15-JUL-1999
15-JUL-1999
  SEQUENCE
           Hypothetical
                     Pfam; PF01062; Worm_family_8;
                                                                        entities requires a license agreement (See or send an email to license@isb-sib.ch).
                                                                                                                between the Swiss Institute of Bioinf
the European Bioinformatics Institute
                                                                                                                         This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation -
                                                                                                                                                                                           SEQUENCE FROM N.A. STRAIN-BRISTOL N2;
                                                                                                                                                                                                                                     Eukaryota; Metazoa;
Rhabditidae; Pelode;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REPEAT
REPEAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REPEAT
REPEAT
                                InterPro;
                                          WormPep; F32B6.9; CE09864.
                                                   EMBL; Z81074; CAB03043.1;
                                                                                                                                                          Submitted (OCT-1996) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                 Basham V.;
                                                                                                                                                                                                                           NCBI_TaxID=6239;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRINTS; PR00319; GPROTEINB.
PRINTS; PR00320; GPROTEINBRPT.
PR051TE; PS00678; WD_REPEATS_1; 3.
PROSITE; PS50082; WD_REPEATS_2; 6.
PROSITE; PS50294; WD_REPEATS_REGION;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  InterPro; IPR001632; -
InterPro; IPR001680; -
Pfam; PF00400; WD40; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EMBL; AF022086; AAB82553.1; HSSP; P04901; IGP2. MGD; MGI:101848; Gnb5.
                                                                                                                                                                                                                                                        Caenorhabditis elegans.
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                                IPR000615;
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11 protein.
                                                                                                                                                                                                                                    Peloderinae;
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                                                                                                                                                                                                                                                                          38, Created)
38, Last sequence update)
38, Last annotation update)
KDA PROTEIN F32B6.9 IN CHROMOSOME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    :; WD repeat;
91 WD
133 WD
181 WD
225 WD
267 WD
311 WD
4.8965
                                                                                                                                                                                                                                             Nematoda; Chromadorea; Rhabditida; Rhabditoidea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 38731 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                        57.9%;
70.0%;
 WW.
                                                                                                                                                                                                                                   Caenorhabditis
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                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 33; DB Pred. No. 20;
                                                                                                                                                                                                                                                                                                                                 PRT;
A66E69A83C78790B CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Multigene
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                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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                                                                                                     There are no rest
                                                                                                                                                                                                                                                                                                                                 A
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                                                                                  http://www.isb-sib.ch/announce/
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                                                                                                                 restrictions
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no
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165

TEKEYQILEA 174

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TRPG_YEAST
ID TRPG_YEAST
AC P00937;
     RROCCOSEE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       identified the known loci URA1, SAC1 and TRP3, and revealed 6 new open reading frames including homologues to the threonine dehydratases, membrane transporters, hydantoinases and the phospholipase A2-activating protein.";
Yeast 10:663-679(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Aebi M., Furter R., Prantl F., Niederberger P., Huetter R.; "Structure and function of the TRP3 gene of Saccharomyces cerevisiae: analysis of transcription, promoter sequence, and sequence coding for
                                                entities requires a license agreement (S or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE-84162082; PubMed-6323449;
Zalkin H., Paluh J.L., van Cleemput M., Moye W.S.
"Nucleotide sequence of Saccharomyces cerevisiae
encoding bifunctional anthranilate synthase: indephosphate synthase.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Saccharomyces cerevisiae (Baker's yeast).

Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Saccharomyces.
                                                                                                 use by non-profit institutions as long
                                                                                                                                                                                                                                                                                                                                                                              ÷
                                                                                                                                                                                                                                                                                                                                                                                                                                            a glutamine amidotransferase.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRAIN-S288C
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01-JUN-1994 (Rel. 29,
15-JUL-1999 (Rel. 38,
ANTHRANILATE SYNTHASE
                                                                                                                                                       This
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=95028164;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Submitted
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=4932;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Zermia M., Horaitis O., Alexandraki D.;
The complete sequencing of a 24.6 kb se
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRAI)
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                                                                                                                                                                                                                                                                                DS-PHOSPHATE = 1-(INDOL-3-YL)GLYCEROL 3-PHOSPHATE + CO(2) + H(2) PATHWAY: FIRST AND FOURTH STEPS IN BIOSYNTHESIS OF TRYPTOPHAN SUBUNIT: TETRAMER OF TWO COMPONENTS I AND TWO COMPONENTS II. INDUCTION: BY TRYPTOPHAN STARVATION AND TWO COMPONENTS II. MISCELLANBOUS: COMPONENT I CATALYZES THE FORMATION OF ANTHRANI INTER TWANTED MINISTELLANBOUS:
                                                                                                                                                                                                                                                                                                                                                         CATALYTIC ACTIVITY: 1-(2-CARBOXYPHENYLAMINO)-1-DEOXY-D-RIBULOSE
5-PHOSPHATE = 1-(INDOL-3-YL)GLYCEROL 3-PHOSPHATE + CO(2) + H(2)(
                                                                                                                                                                                                                                                                                                                                                                                              CATALYTIC ACTIVITY: CHORISMATE + L-GLUTAMINE = PYRUVATE + L-GLUTAMATE.
                                                                                                                                                                                                       MISCELLANEOUS: YEAST COMPONENT II CARBOXYL END ALSO INDOLE-3-GLYCEROL PHOSPHATE SYNTHASE ACTIVITY. SIMILARITY: THE GATASE DOMAIN BELONGS TO TYPE-1 GLUT
                                                                                                                                                                                                                                                       USING AMMONIA RATHER THAN GLUTAMINE, WHEREAS GLUTAMINE AMIDOTRANSFERASE ACTIVITY.
                                                                                                                                                                                       AMIDOTRANSFERASES.
                                                                                                                                SWISS-PROT entry is copyright. It is producen the Swiss Institute of Bioinformatics
; K01386;
; X75951;
                                                                                                                                                                                                                                                                                                                                                                                                                              Genet. 8:165-172(1984)
                                                                                 and this statement is not removed
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Last annotation update)
COMPONENT II (EC 4.1.3.27) [INCLUDES:
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                                                            (See http://www.isb-sib.ch/announce/
                                                                                                There are no rest
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4.1.1.48)
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EMBL;
PIR; S
                                                                                                                       McMahon A., Geertman R., Sabban E.L.;
McMahon A., Geertman R., Sabban E.L.;
"Rat dopamine beta-hydroxylase: molecular cloning a
"Rat dopamine beta-hydroxylase: molecular cloning a
                                                                                                                                                                                                                                                    01-FEB-1995 (Rel. 31, Created)
01-FEB-1995 (Rel. 31, Last sequence update)
01-OCT-2000 (Rel. 40, Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRINTS; PR00096; GATASE.
PRINTS; PR00097; ANTSNITHASEII.
PROSITE; PS00442; GATASE_TYPE_I; 1.
PROSITE; PS00614; IGPS; 1.
Tryptophan biosynthesis; Transferas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      InterPro;
InterPro;
                                              -1- COFACTOR: PQQ, COPPER, AND ASCORBATI
-1- PATHWAY: CATECHOLAMINE BIOSYNTHESIS
-1- SUBUNIT: HOMOTETRAMER.
                                                                          SEQUENCE FROM N.A. MEDLINE-90219125; PubMed-2325165;
                                                                                                                                                                                           Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                               Rattus norvegicus (Rat).
                                                                                                                                                                                                                           DBH
                                                                                                                                                                                                                                   HYDROXYLASE)
                                                                                                                                                                                                                                            01-0CT-2000 (Rel. 40, Last annotation update)
DOPAMINE BETA-MONOOXYGENASE PRECURSOR (EC 1.14.17.1) (DOPAMINE BETA-
                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Glutamine amidotransferase.
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                                                                                                                reserpine.";
                                                                                                                                                                                  NCBI_TaxID=10116
                                                                                                                                                                                                                                                                                                                                             348
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        FORM IN CHROMAFFIN GRANULES (THE MEMBRANE-BOUND FORM MAY BE ANCHORED BY AN UNCLEAVED SIGNAL SEQUENCE).
TISSUE SPECIFICITY: CHROMAFFIN GRANULES OF THE ADRENAL MEDULLA
                                     SUBCELLULAR LOCATION: EXISTS AS BOTH
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$44327; $44327.
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M36300;
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IPR002385;
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AAA34450.1;
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                                                                                                                                                                                                                                                                                            STANDARD;
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Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                          57.98;
77.88;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Transferase; Lyase; Multifunctional enzyme;
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                                                               AND ASCORBATE.
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GATASE (BY SIMILAR
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Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                             PRT;
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SYMPATHETIC NERVOUS
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e mRNA
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                                     AND
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; Murinae; Rat
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                             MEMBRANE-BOUND
RM MAY BE
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This

MONOOXYGENASE FAMILY

SIMILARITY: BELONGS TO THE COPPER TYPE II, ASCORBATE-DEPENDENT

SWISS-PROT entry is copyright. It is produced through a collaboration sen the Swiss Institute of Bioinformatics and the EMBL outstation .

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Best Local Similarity
Matches 6; Conser
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CJBA_BACUH STANDARD,
CR6170;
O86170;
O1-OCT-2000 (Rel. 40, Created)
O1-OCT-2000 (Rel. 40, Last sequence update)
O1-OCT-2000 (Rel. 40, Last annotation update)
PESTICIDIAL CRYSTAL PROTEIN CRY199A (INSECTICIDAL DELTA-ENDOTOXIN
PESTICIDIAL CRYSTALINE ENTOMOCIDAL PROTOXIN) (78 KDA CRYSTAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CARBOHYD
CARBOHYD
SEQUENCE
                                                                                     SEQUENCE FROM N.A.

MEDLINE=98369730; PubMed=9704107;

Hwang S.H., Saitoh H., Mizuki E., Higuchi K., Ohba M.;

Hwang S.H., Saitoh H., Mizuki E., Higuchi K., Ohba M.;

"A novel class of mosquitocidal delta-endotoxin, Cry19B, encoded by a Bacillus thuringiensis serovar higo gene.";

Bacillus thuringiensis serovar higo gene.";

Syst. Appl. Microbiol. 21:179-184(1998).

-i- FUNCTION: PROMOTES COLLOIDOSMOTIC LYSIS BY BINDING TO THE MIDGUT
                                                                                                                                                                                                                                                                                           Bacteria; Firmicutes; Bacillus/Clostridium group; Bacillus/Staphylococcus group; Bacillus: NCBI_TaxID=132266;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CARBOHYD
CARBOHYD
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                                                                                                                                                                                                                                                                                                                                                                         Plasmid.
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PROSITE; PS00084; CU2_MONOOXYGENASE_1;

PROSITE; PS00085; CU2_MONOOXYGENASE_2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    the European Bioinformatics Institute. The use by non-profit institutions as long modified and this statement is not removed
                                                                                                                                                                                                                                                                                                                                                                                             Bacillus thuringiensis
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InterPro; IPR001178; -. Pfam; PF00555; endotoxin; 1. Toxin; Sporulation; Plasmid. SEQUENCE 682 AA; 78490 MW;
                                                                                                                                                                                                                    use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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ALIGNMENTS

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J. Clin. Invest. 90:2102-2109(1992).
                                                                                                                                                                                                                                                                                                                                                          MEDLINE=93055454; PubMed=1430234;
Porges A.J., Redecha P.B., Doebele R.,
Kimberly R.P.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ₹
                                                                                                                                                                   Length 374;
                                                                                                                                                                                                                                                                                                                                                                             Salmon J.E.,
                                                                                                                                           Indels
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                                                                                                                                                                                                                                                                    Matches
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Best Local
Eukaryota; Viridiplantae; Embryophyta; Magnoliophyta; eudicotyledons; core eud Brassicales; Brassicaceae. ***
                                                                                                                       01-OCT-2000 (TrEMBLrel.
01-OCT-2000 (TrEMBLrel.
01-OCT-2000 (TrEMBLrel.
                                                                                                                 T3F20.14 PROTEIN.
                                                                                                                                                        Q9LPH0
                                                                   NCBI_TaxID=3702
                                                                                                           T3F20.14
                                                                                                                                                 Q9LPH0
                                                                                                                                                                                                                                                                                                                 EMBL;
EMBL;
EMBL;
                                                                                                                                                                                                                                                                        INTERPRO; IPRO03006; PFAM; PF00047; ig; 3; PRODOM; PD002534; -;
                                                                                                                                                                                                                                                                                                                                       EMBL;
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                                                                                                                                                                                                                                                                                                                                                      EMBL;
                                                                                                                                                                                                                                                                                                                                                                     EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE OF 1-224 FROM N.A. MEDLINE-93018827; PubMed-1402657; Benech P.D., Sastry K., Iyer R.R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-FEB-1997 (TrEMBLrel. 02, Created)
01-FEB-1997 (TrEMBLrel. 02, Last sequence update)
01-OCT-2000 (TrEMBLrel. 15, Last annotation update)
FC GAMMA RECEPTOR TYPE I (FC GAMMA RIB=FC GAMMA RECEPTOR).
CD64 OR FC-GAMMA>RIB.
                                                                                                                                                                                                                                                                                                                                                                                    J. Exp. Med. 176:1115-
EMBL; M91555; AAA58414
                                                                                                                                                                                                                                                                                                                                                                                                     structure
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TISSUE-BLOOD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                            gamma receptor gene
                                                                                                                                                                                                                                                                                                                                                                                                                  *Definition of interferon
                                                                                                                                                                                                                                                                                                                                                                                                                            Ezekowitz R.A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ezekowitz A.B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Benech P.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              092495;
01-FEB-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EQUENCE FROM N.A.
                                                                                                                                                                                                                                   Local
                                                                                                                                                                                                             TSSEYQILTARR
                                                                                                                                                                                              TSSEYQILTARR
                                                                                                                                                                                                                                                                                                 M91554;
S45709;
S45707;
S45708;
S45704;
S45705;
P12319;
                                                                                                                                                                                                                            l Similarity
12; Conser
                                                                                                                                                                                                                            ilarity 100.
Conservative
                                                                                                                                                                                                                                                                   375
                                                                                                                                                      PRELIMINARY;
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                                                                                                                                                                                                                                                                                                         AAD13842.1;
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AAA58414.1;
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                                                                                                                     Last sequence update)
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Pred.
                                                                                                                                                       PRT;
                                                                                                                                                                                                                                                                                                                                                                                                           RIb)
                                                                               core eudicots;
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                                                                                                                                                                                                                                                                 A84D464C70DD0F91 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R.R.,
                                                                                                                                                                                                                           Mismatches
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                                                                              Tracheophyta; s
dicots; Rosidae;
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                                                                              Spermatophyta; e; eurosids II;
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Best Local S
Matches 7
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01-OCT-2000
01-OCT-2000
SEQUENCE FROM N.A.

Sherburne C.K., Lawley T.D., Gilmour M.W.,
Grotbeck E., Rose D.J., Taylor D.E.;
"The complete DNA sequence and analysis of
from Salmonella typhi that is temperature s
Nucleic Acids Res. 28:2177-2186(2000).
EMBL; AR250878; AAF69908.1; -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ecker J.R., rederspiel N.A., Theologis A.;
"The sequence of BAC T3F20 from Arabidopsis tl Submitted (JUN-2000) to the EMBL/GenBank/DDBJ EMBL; ACOL8748; AAF78435.1; -.
                                                                                             Plasmid R27.
Bacteria; Proteobacteria;
Salmonella.
                                                                                                                         Salmonella
                                                                                  NCBI_TaxID=601;
                                                                                                                                   R0070
                                                                                                                                           HYPOTHETICAL
                                                                                                                                                                                Q9L5R1;
                                                                                                                                                                                          Q9L5R1
                                                                                                                                                                                                                                                                                                                                       "Wan Y., Li X., Wang A., Zhang G.;
"Molecular cloning and identification of high affinity Fc receptor for bovine Igg Vet. Immunol. Immunopathol. 75:151-159(20 EMBL; AF162866; AAF80477.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                        Bos taurus (Bovine).

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eute
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora;
Bovidae; Bovinae; Bos.

NCBI_TaxID=9913;
                                                                                                                                                                                                                                                                                                                          SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-0CT-2000 (TrEMBLrel. FC GAMMA RECEPTOR I.
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01-OCT-2000
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                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE
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                                                                                                                                                                                                                                                        1 TSSEYQILTARR
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                                                                                                                                                                                        PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                         AA;
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                                                                                                                                                                                                                                                                                                                       39608 MW;
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58
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15,
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Last
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Pred. No.
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Pred.
                                                                                                                                                                                       PRT;
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                                                                                                   subdivision; Enterobacteriaceae;
                                                                                                                                                                                                                                                                         Mismatches
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14;
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                          f R27, a large sensitive for
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                                    R27,
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                                                     Blattner F.R.,
                                                                                                                                                                                                                                                                                           6; Length 349;
                                                                                                                                                  update)
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                                                                                                                                                                                                                                                                         3,
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                          IncHI plasmid
transfer.";
                                                                                                                                                                                                                                                                                                                                                                                                                           Bovoidea;
                                                     Burland
                                                                                                                                                                                                                                                                                                                                                                    encoding
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SEQUENCE

791

89568 MW;

46D6F5751A1C7D67 CRC64;

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RESULT
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Best Local
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Best Local Similarity
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01-MAY-2000
01-MAY-2000
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                                                                                                                                                                                                                                                                                   Q9VWA3;
                                                                                                     Eukaryota; Metazoa; I
Pterygota; Neoptera;
                                                                                                                                                                              CG13814
                                                                                                                                                                                                CG13814 PROTEIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PFAM; PF01377; Thioester_dehyd; 2.

PROSITE; PS00606; B_KETOACYL_SYNTHASE; UNKNOWN_1.

PROSITE; PS00881; PROTEIN_SPLICING; UNKNOWN_1.

PRODOM; PD013327; -; 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
STRAIN-SCRC-2738;
MEDLINE-97419510; PubMed-9274025;
Takeyama H., Takeda D., Yazawa K., Y
Takeyama hours of the elcosapentaenoic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-JAN-1998 (TrEMBLrel. 05, Cre 01-JAN-1998 (TrEMBLrel. 05, Las 01-OCT-2000 (TrEMBLrel. 15, Las HYPOTHETICAL 216.9 KDA PROTEIN. Shewanella sp. SCRC-2738.
SEQUENCE FROM N.A. STRAIN-BERKELEY;
                                                                                                                                                     Drosophila melanogaster
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hypothetical protein. SEQUENCE 2004 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PFAM; PF00109; ketoacyl-synt; PFAM; PF01377; Thioester_dehyd
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Microbiology 143:0-0(0).
EMBL; U73935; AABB1125.1;
HSSP; P18391; 1MKA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bacteria; Proteobacteria; gamma subdivision; Alteromonadaceae;
                                                                                        Ephydroidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Shewanella
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                                                                 NCBI_TaxID=7227;
                                                                                                                                                                                                                                                                                                                                                                                                                    905
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=53560;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity 72.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 TSSEYQILTAR 11
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                                                                                                                                                                                                                                                                                                          PRELIMINARY;
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                                                                               Arthropoda; Tracheata; Hexapoda; Insecta;
; Endopterygota; Diptera; Brachycera; Muscomorpha;
hilldae; Drosophila.
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72.7%;
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c acid synthesis gene cluster fro
ine cyanobacterium, Synechococcus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 36;
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Best Local
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Q18337;
Q1-NOV-1996 (T
Q1-NOV-1996 (T
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COSMID C31H1.
                                                       SEQUENCE FROM N.A.
MEDLINE=94150718; PubMed-7906398;
Wilson R., Ainscough R., Anderson
Bonfield J., Burton J., Connell M
                                                                                                                                                                                                Caenorhabditis elegans.
Eukaryota; Metazoa; Nemato
Rhabditidae; Peloderinae;
NCBI_TaxID=6239;
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EMBL; AE003514; AAR49045.1; -.

FLYBASE; FB9G003560; CG13814.

SEQUENCE 104 AA; 11491 MW; 2620F15C7CBFB431 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Shue B.C., Siden-Kiamos I., Simpson M., Skupski M.P., Smith T., Spier E., Spradling A.C., Stapleton M., Strong R., Sun E., Svirskas R., Tector C., Turner R., Venter E., Wang A.H., Wang X., Wang Z.-Y., Wassarman D.A., Weinstock G.M., Weissenbach J., Williams S.M., Woodage T., Worley K.C., Wu D., Yang S., Yao Q.A., Ye J., Yeh R.-F., Zaveri J.S., Zhan M., Zhang G., Zhao Q., Zheng
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Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottler P.,
Burtis K.C., Busam D.A., Butler H., Cadieu E., Center A., Chandra I.,
Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
de Pablos B., Delcher A., Deng Z., Mays.A.D., Dew I., Dietz S.M.,
Dodson K., Doup L.E., Downes M., Dugan-Roccha S., Dunkov B.C., Dunn P.
                                                                                                                                                                                                                                                                                                                           С31Н1.2
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                           Craxton M.,
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Ainscough R., Anderson K., Bayn
, Burton J., Connell M., Copsey
Dear S., Du Z., Durbin R., Fay
Green P., Hawkins T., Hillier
                                                                                                                                                                                                                                                                                                                                                                                                                                         (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                     (TrEMBLrel.
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                                                                                                                                                                                                                                                                       Nematoda;
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Caenorhabditis.
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                                                    Copsey T., Cooper J.,
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                           Favello A., Fulton
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ri V., Reese M.G.,
                              Fulton
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X., Smith H.O.,
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                                                             Coulson A.,
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Query Match
Best Local Similarity
7; Conserv
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A Addams M.D., Celniker S.E., Li P.W., Hoskins R.A., Galle R.F.,

A George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N.,

A Sutron G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X.,

A Brandon R.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfeiffer B.D.,

A Wan K.H., Doyle C., Baxter E.G., Helt G., Nelson C.R., Miklos G.L.G.,

A Wan K.H., Dayle C., Baxter E.G., Helt G., Nelson C.R., Miklos G.L.G.,

A Wan K.H., Dayle C., Baxendale J., Bayraktaroglu L., Beasley E.M.,

RA Ballew R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M.,

RB Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S.,

RA Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P.,

RA Burtis K.C., Busam D.A., Butler H., Cadieu E., Center A., Chandra I.,

RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,

RA de Pablos B., Delcher A., Deng Z., Mays A.D., Dew I., Dietz S.M.,

RA de Pablos B., Delcher A., Deng Z., Mays A.D., Dew II., Dietz S.M.,
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              Dodson K., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P., Durbin K.J., Evangelista C.C., Ferraz C., Ferriera S., Fleischmann W., Fosler C., Gabrielian A.E., Garg N.S., Gelbart W.M., Glasser K., Glodek A., Gong F., Gorrell J.H., Gu Z., Guan P., Harris M., Harvey D., Heiman T.J., Hernandez J.R., Houck J., Hostin D., Houston K.A., Howland T.J., Wei M.-H., Ibegwam C., Jalali M., Kalush F., Karpen G.H., Ke Z., Kennison J.A., Ketchum K.A., Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z., Lasg Y., Lin X., Lasko P., Lei Y., Levitsky A.A., Li J., Li Z., Liang Y., Lin X.,
      Lasko P.
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01-MAY-2000
01-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Drosophila melanogaster (Fruit fly).

Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;

Eukaryota; Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;

Ephydroidea; Drosophilidae; Drosophila.
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INTERRENC, IPRO00215; -
PROSITE: PS00284; SERPIN; UNKNOWN_1.
SEQUENCE 197 AA; 22503 MW; 00A5A7EB028E20F4 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=20196006;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Jones M., Kershaw J., Kirsten J., Laister N., Latreille P., Lightning J., Lloyd C., Mcmurray A., MortLore B., O'Callaghan M., Parsons J., Percy C., Rifken L., Roopra A., Saunders D., Shownkeen Smaldon N., Smith A., Sonnhammer E., Staden R., Sulston J., Thierry-Mieg J., Thomas K., Vaudin M., Vaughan K., Waterston R., Watson A., Weinstock L., Wilkinson-Sproat J., Wohldman P., "2.2 Mb of contiguous nucleotide sequence from chromosome III of C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Waterston
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     PubMed=10731132;
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58.3%;
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Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
McLeod M.P.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  35;
No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     553
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
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RESULT 11 .Q9SZR4 ID Q9SZR4

PRELIMINARY;

PRT;

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41

TRSEFQPMTAR 51

1 TSSEYQILTAR 11

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                                                  Query Match
Best Local
            Matches
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                 EMBL; AF104240; AAD21011.1; HSSP; P20783; 1B8K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Jansen H.T., Lehman M.N., Stevens P.J.;
"Golden Hamster Neurotrophin and Neurotrophin
Submitted (NOV-1998) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.,
Jansen H.T., Lehman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mesocricetus auratus (Golden hamster).
Eukaryota; Metazoa; Chordata; Craniata
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Wang Z.-Y., Wassarman D.A., Weinstock G.M., Weissenbach J., Williams S.M., Woodage T., Worley K.C., Wu D., Yang S., Yao Q.A., Ye J., Yeh R.-F., Zaveri J.S., Zhang G., Zhao Q., Zheng L., Zheng X.H., Zhong F.N., Zhong W., Zhou X., Zhu X., Zhu X., Smith H.O., Gibbs R.A., Myers E.W., Rubin G.M., Venter J.C., "The genome sequence of Drosophila melanogaster."; science 287:2185-2195(2000).
                                                                                                                                                                                                                   SEQUENCE
                                                                                                                                                                                                                                                                                                                                     PFAM; PF00243; NGF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=10036;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mesocricetus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INTERPRO; IPR000822; -. PFAM; PF00096; zf-C2H2; 2. PFAM; PF00651; BTB; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nelson D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M., Palazzolo M., Pittman G.S., Pan S., Pollard J., Puri V., Reese M Reinert K., Remington K., Saunders R.D.C., Scheeler F., Shen H., Shue B.C., Siden-Kiamos I., Simpson M., Skupski M.P., Smith T., Spier E., Spradling A.C., Stapleton M., Strong R., Sun E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EMBL; AE003435; AAF46100.1; -. FLYBASE; FBgn0029822; CG12236.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Svirskas R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Merkulov G., Milshina N.V., Mobarry C. Mount S.M., Moy M., Murphy B., Murphy
                                                                                                                                                                                                                                                                                                                                                                           INTERPRO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PROSITE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   363
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2 SSEYQILT 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SSEYEILT 370
        Similarity 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PS00028; ZINC_FINGER_C2H2; 2.
553 AA; 60801 MW; EB3019D
                                                                                                                                                                                                                                                                                                                                                                       783; 1B8K.
IPR002072; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Spradling A.C., Stapleton M., Strong R., Sun E., R., Tector C., Turner R., Venter E., Wang A.H., Wang X.,
                                                                                                                                                                                                               129
129 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    IPR000210;
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                                                                                                                                                                                                               14890 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Chordata; Craniata; Vertebrata; Euteleostomi; Rodentia; Sciurognathi; Muridae; Cricetinae;
                                              59.6%;
63.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             61.4%;
2;
        Score 34; DB
Pred. No. 18;
2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 35; DB Pred. No. 53; 1; Mismatches
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                                                                                                                                                                                                               43C825E6651EFDF3 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EB3019D282B19092 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              update)
                                                                                    11;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Receptor cDNA.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0,
                                                                                    Length 129;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Moshrefi A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0,
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Query Match Best Local (

59.6%;

Score 34; Pred. No.

DB

Length 863;

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В
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Best Local
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01-MAY-2000
01-MAY-2000
01-OCT-2000
                                                                                                                                                                                                                                  Q9P5K7
Q9P5K7;
           German Neurospora genome project;
Submitted (MAY-2000) to the EMBL/
EMBL; AL356172; CAB91690.1;
                                                                       Schulte U., Aign V.,
Nyakatura G., Mewes
Submitted (MAY-2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EU Arabidopsis sequencing pr
Submitted (MAR-2000) to the
EMBL; ALOSO352: CAB43668 1;
EMBL; AL161575; CAB79751.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bevan M., Zimmermann W.,
Mewes H.W., Mayer K.F.X.,
Submitted (MAY-1999) to t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Arabidopsis thaliana (Mouse-ear cress).
Eukaryota; Viridiplantae; Embryophyta; Trachec
Magnoliophyta; eudicotyledons; core eudicots;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-CT-2000 (TrEMBLrel. 15, Last annotation update)
HYPOTHETICAL 32.0 KDA PROTEIN.
F27B13.170 OR AT4G29930.
                                                                                                                                 Eukaryota; Fungi;
NCBI_TaxID=5141;
                                                                                                                                                                                              01-OCT-2000
01-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Brassicales; Brassicaceae; Arabidopsis.
NCBI_TaxID=3702;
 SEQUENCE
                                                SEQUENCE FROM N.A.
                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                               Neurospora crassa.
Eukaryota; Fungi;
                                                                                                                                                                    B23L21.220
                                                                                                                                                                                   RELATED TO GLUCAN
                                                                                                                                                                                                                                                                                                                                                                                                                    Hypothetical protein.
SEQUENCE 277 AA; 31987 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Submitted
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Submitted
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ammermann
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                                                                                                                                                                                                                                                                                                           257
                                                                                                                                                                                                                                                                                                                                                                                                                                           PROSITE; PS00038; HELIX_LOOP_HELIX; UNKNOWN_1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NTERPRO;
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                                                                                                                                                                                                                                                                      12
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7; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mewes H.W., Lemcke K., Mayer K.F.X.; (MAR-2000) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    IPR003015; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IPR001092;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              opsis sequencing project; (MAR-1999) to the EMBL/GenBank/DDBJ
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863 AA;
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(TrEMBLrel. 15, Last sequence update)
(TrEMBLrel. 15, Last sequence update)
(TrEMBLrel. 15, Last annotation update)
                                                                                                                                                                                                                                               PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                         HLH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Grueneisen A.,
                                                                                                                                               Ascomycota;
                                                                      , Hoheisel J., Brandt P., I
H.W., Mannhaupt G.;
) to the EMBL/GenBank/DDBJ
 98928 MW;
                                                                                                                                                                                                                                                                                                                                                                      59.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              W., Grueneisen A., F.X., Schueller C.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                project;
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                        EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                         2;
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Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Wambutt R.,
                                                                                                                                                                                                                                               PRT;
                                                                                                                                               Sordariales;
 EC64BADA28708040 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                     2E85E0FE320F073F CRC64;
                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                      No 4
                                                                                                                                                                                                                                             863
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tracheophyta; Spermatophyta; dicots; Rosidae; eurosids II;
                                                                                                                                                                                                                                                                                                                                                                                  В
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                                                                                                                                               Sordariaceae;
                                                                                              Fartmann B.,
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••
                        databases
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                                                                                                                                               Neurospora
                                                                                                 Holland
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01-MAY-2000
01-MAY-2000
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Q9LTX7;
Q1-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                              EMBL; ABO
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                               features of the region clones.";
DNA Res. 7:31-63(2000)
                                                                               "Novel dendritic kinesin sorting identified by different targeting of two related kinesins: KIF21A and KIF21B.";
J. Cell Biol. 145:469-479/1999
                                                    EMBL; AF202893; AAF17084
HSSP; P17119; 3KAR.
MGD; MGI:109234; Kif21b.
                                                                                                           MEDLINE=99242603; PubMed=10225949; Marszalek J.R., Weiner J.A., Farlo
                                                                                                                                                         Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                            Mus musculus
                                                                                                                                                                                                                                            Q9QXL1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A. STRAIN-COLUMBIA;
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01-OCT-2000
               PFAM; PF00225; kinesin; 1. PFAM; PF00400; WD40; 7.
                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                               NCBI_TaxID=10090;
                                                                                                                                                                                       KIF21B.
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                                   INTERPRO; IPRO01680; -.
INTERPRO; IPRO01752; -.
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NCE 1080
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7; Conserv
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6; Conser
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etazoa; Chordata;
theria; Rodentia;
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                                                                       AAF17084.1;
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AA; 121430
                                                                                                                                                                                                                                                                                                                                                                                                                    regions
        KINESINHEAVY
KINESIN_MOTOR_DOMAIN1; UNKNOWN_1
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Pred. No. 1.8e
1; Mismatches
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Sciurognathi; Muridae
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Murinae; Mus
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Best Local Similarity 54.5
Matches 6; Conservative
                                                             Query Match 57.9%;
Best Local Similarity 70.0%;
Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PROSITE; PS00678; WD_REPEATS; UNKNOWN_1.
PROSITE; PS50067; KINESIN_MOTOR_DOMAIN2; 1.
SEQUENCE 1668 AA; 186162 MW; 37145B2D867C9A57 CRC64;
                                                                                                                                                                                                                                                                                                                                                   Q9NPW7 PRELIMINARY; PKT; 2/1 An.
Q9NPW7;
Q1-QCT-2000 (TrEMBLrel. 15, Created)
Q1-QCT-2000 (TrEMBLrel. 15, Last sequence update)
Q1-QCT-2000 (TrEMBLrel. 15, Last annotation update)
HYPOTHETICAL 30.8 KDA PROTEIN (FRAGMENT)
DKFZP762M186.
DKFZP762M186.
UNMAN)
                                                                                                                                                                                               SEQUENCE FROM N.A.

TISSUB-MELANOMA (MEWO CELL LINE);
Bloecker H., Boecher M., Brandt P., Mewes H.W., Weil B., Wiemann S.;
Submitted (JUN-2000) to the EMBL/GenBank/DDBJ databases.

EMBL; AL359590; CAB94876.1; -.
                                                                                                                                                                                                                                                                                                             Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                Hypothetical protein:
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SEQUENCE 271 AA; 3
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259 TGTEYETLTAK 269
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1 TSSEYQILTA 10
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117 TRAEYQILLA 126
                                                                                                                                                271 AA; 30764 MW;
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Search completed: June 4, 2001, 12:23:27 Job time: 590 sec

GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.

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Result
No.
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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Maximum DB
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Match
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Gapop 10.0 , Gapext 0.5
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R22550
R91438
W00859
Y96134
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R91439
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Y33183
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             Human macrophage-s
Human FCRI (cDNA c
Human FCRI (cDNA c
Human macrophage-s
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Human EcRI (cDNA c
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high affinit
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	828	789	789	764	738	738	517	398	333	272	4412	761	582	561	1018	1018	410	234	234	150	1728	1447	1447	1447	1251	1242	1139	1125	1018	377	374	374	374	399
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21 TOWNS 100	R13905	W68007	R62488	W69679	Y59302	Y29639	Y32390	Y96307	в12313	R92254	Y53666	R92255	R92256	R91065	R87028	R63759	R12428	Y87334	W74886	Y02365	R13144	в50693	Y33498	R68553	W37778	W52287	W37779	W52288	W06485	W86195	W97834	W97833	4	B43683
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-	Rabbit ATHERO-ELAM	Human soluble vasc	Truncated KLD sVEG	KDR protein sequen	Human VEGF recepto	Human vascular end	Herpesvirus entry		Human secreted pro	e11	ce gi/	cell	Neural cell adhesi	Rat interleukin-1	Human contactin.		Hybrid Fc(gamma)RI	signal p	Human secreted pro	Polypeptide identi	Deleted in Colorec	•	8	ã			Rattus norvegicus	Rattus norvegicus	Rat contactin liga	Human Fc receptor		FC	Fc rece	Human cancer assoc

ALIGNMENTS

RESULT W60558 08-OCT-1996; 07-OCT-1997; WO9815833-A1. Synthetic. cell transformation; mutation; anti; oncogene. Extracellular domain; CD64; identification; antibody; immunohistochemical; immunofluorescent analysis; dete Oligopeptide from extracellular domain of CD64 18-AUG-1998 W60558; W60558 standard; peptide; 12 (UYUT-) RIJKSUNIV UTRECHT 16-APR-1998. (first entry) 96EP-0202791. 97WO-NL00557 immunofluorescent analysis; detection; ₽ .

WPI; 1998-240964/21. De Kruif CA, Logtenberg T;

XX PA

Identifying peptide(s) binding specifically to protein target - by expressing on phage surface and testing for binding to immobilised oligo:peptide derived from the target, useful for, e.g. identifying specific antibodies e.g. identifying

Example 1; Page 29;

40pp; English

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RESULT
Y33183
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Best Local Similarity 100
Matches 12; Conservative
                     Baell JB, E
Maxwell KF,
Powell MS;
                                                                                                                                                                      11-SEP-1998;
06-FEB-1998;
                                                                                                                                                                                                                                                              04-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              anti-inflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    drug design; therapy; inhibitor;
tissue damage; hypersensitivity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 atomic coordinate;
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                                                                                                                (ILEX-) ILEXUS PTY LTD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Fc receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human src-gammaRI protein fragment.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic oligopeptides W60537-61 are derived from the extracellular domain of CD64. They were synthesised on 25 polyethylene rods as adjacent, non-overlapping sets. Peptide specific phages were isolated by affinity binding to the CD64-covered rods. The specification describes a method for the identification of a peptide able to bind specifically to a target protein. The method comprises displaying the peptide on the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         by mutation in (anti)oncogenes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            selected peptides can be isolated and used for recombinant production of antibodies. Selected antibodies are used, e.g. for immunohistochemical or immunofluorescent analysis, and also to detect cell transformation caused
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               nflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 EDSGLYQCEAAT 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ν
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             of the same protein,
         Epa V, var. F, McCarthy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      e; hypersensitivity; inf
modulator; Fc-gammaRIa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Fc-gammaRIIa; human; FCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                   98US-0099994.
98US-0073972.
                                                                                                                                                                                                                                                           99WO-IB00367.
                                                Garrett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nmaRIIa; human; FCR; model; three-dimension; 3 bloactive compound design; computer-assisted;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              immunoprotective;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%;
                        TD,
                     McKenzie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   261 AA:
                                                TPJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 64;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                  Hogarth PM,
zie IFC, Pie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fc-gammaR; Fc-epsilonR; IgG; IgE; inflammatory cell recruitment; RIa; immune function regulation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sfc-gammaRI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.00033;
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                  Pietersz GA;
                                                Matthews
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
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                                                BR;
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Three-dimensional structures and models of Fc receptors, useful in computer-assisted drug design $% \left(1\right) =\left\{ 1\right\}$

This amino acid sequence was predicted from the cDNA sequence of CDNA clone p98/X2. It differs from the sequence predicted from clone p185 (see Q21178) at position 25, p135 encodes a Ser residu while clone p98/X2 predicts a Thr residue. At position 58, p90

Ser residue 58, p90

Example 10;

Page 94a; 160pp; English.

New CD53 cell surface antigen and DNA encoding it - for immuno-therapy and diagnosis of haematopoietic neoplasms, etc.

WPI; 1999-539978/45

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E PD PN XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   that inhibit the activity of Fc-gammaR or Fc-epsilonR can be used to reduce IgG- or IgE-mediated tissue damage, e.g. IgG- or IgE-mediated hypersensitivity; recruitment of inflammatory cells or release of inflammatory modulators. The therapeutic compositions can also be used to treat disease in animals. The knowledge of the 3-D structure of Fc-gamma RIA and models of other FCR provides a means for designing and producing compounds that regulate immune function and inflammation in an animal, including humans (i.e. structure based drug design). For example, chemical compounds can be designed to block binding of immunoglobulin to
                                                                                                                                                                         WPI; 1992-056864/07
                                                                                                                                                                                                                                                                                                                                                                                                                                 Rapid immunoselection cloning technique; cell surface antigen immunodiagnosis; high affinity receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human macrophage-specific FcRI receptor encoded by clone p98.
                                                                                                                                                         N-PSDB; Q21179.
                                                                                                                                                                                                                                         (GEHO-) GEN HOSPITAL CORP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R22549 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        an PC receptor protein using various computer programs and models. The products of the invention have anti-inflammatory and immunoprotective activity. This sequence represents the human sFC-gammaRI protein fragment described in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the FcR can be used to design bioactive chemical compounds, e.g. oligonucleotides, peptides, peptidomimetics and small organic molecules, by computer-assisted methods of drug design. Therapeutic compositions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention describes a novel model of an Fc receptor (FCR) protein representing a three-dimensional (3-D) structure that substantially conforms to the specified atomic coordinates. Computer model images of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 11; Page 316-317; 326pp; English
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                                                                                                                                                                                                         Aruffo A,
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Pred. No. 0.098;
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RESULT
R91439
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Best Local S
Matches 11
                                                                                                                                                                                                                                                                                                                                                                                    13-JUL-1989;
13-JUL-1990;
18-OCT-1993;
                 The amino acid sequence (R91439) of human FCRI was detd. from a cDNA library using an immunoselection cloning method. FCRI is a high affinity receptor for the Fc portion of IgG, normally located on cell surfaces of macrophages. 2 Other cDNA clones, pl35 (T14717) and p90 (T14719), coded for a variants of the FCRI sequence; the C-terminal sequence
                                                                                                                                                                                           Cloning of cDNA encoding cell surface antigen of diagnostic and therapeutic proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                             01-DEC-1992;
25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cys residues distributed among three C2 set Ig-related domains. A fusion protein of FCRI and a receptor ligand will be helpful to increase the potency of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sequence, a single and a short, highly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (see Q21180) predicts Val and p98/X2 predicts Leu. Sequences perdicted from all 3 clones show the typical features of a type I integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane-spanning domair
                                                                                                                                                                                                                                                    N-PSDB;
                                                                                                                                                                                                                                                                                                           Aruffo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          09-APR-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cell surface
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R91439;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R91439 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     and a short, highly charged cytoplasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and s
                                                                                                                                                   Example 10; Column 55-56; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-FEB-1988;
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                                                                                                                                                                                                                                                                                                                                               (GEHO)
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253 edsglywceaat 264
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DB; T14718.
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11; Conservative
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                                                                                                                                                                                                                                                                                                                                               HOSPITAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antigen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                  92US-0983647.
88US-0160416.
89US-0379076.
90US-0553759.
93US-0139273.
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                                                                                                                                                                                                                                                                                                                                               CORP.
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ector; FcRI; Fc receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "amino acid
translated
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Pred. No.
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1 product"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    in clone
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those of the pl35
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                                                                                                                                                                                                            useful for isolation
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                                                                                                    Misc-difference
                                                                                                                                      Misc-difference
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                 Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                        amyloidosis; multiple sclerosis; parasitic disease; Fc receptor
                                                                                                                                                                                                                                                                                                                                                                                  panning; immunod
immune.disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                  Human macrophage-specific FcRI.
                                                                                                                                                                                                                                                                                                                                                                                                                                     19-DEC-2000
                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                 Macrophage;
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11; Conser
                                                                                                                                                                                                                                                                                                                                                                                  ge; FCRI; cell surface antigen; human; immunoselect
immunodiagnosis; diagnosis; immunotherapy; gene th
isorder; infection; asthma; immune-complex disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             344 AA;
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9 117
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116
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213
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                  140
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                                                                                                                             TTC"
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                                                                                                                                                               GAA"
                                                                                                                                                                                ÇAA"
                                                                                                                                                                                                  CAG
                                                                                                                                                                                                                   CCC"
                                                                                                                                                                                                                                    TCC.
                                                                                                                                                                                                                                                                     ACC"
                                                                                                                                                                                                                                                                                      cgg"
                                                                                                                                                                                                                                                                                                       ACC"
                                                                                                             ACC '
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     W00859). A helpful in
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A fusion protein in increasing the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 344;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                          therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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immune-complex disease, amyloidosis, parasitic diseases or multiple sclerosis. FcRI is a high affinity receptor for the Fc portion of IgG, normally located on the cell surfaces of macrophages. The ability to interfere with such bonding, or to cause it to occur on surfaces other than macrophages, is useful in therapy. A fusion protein of FcRI and a receptor ligand will be helpful to increase
                                   surfaces other than macrophages, protein of FCRI and a receptor 1 the potencies of antibodies in t
                                                                                                                                                                                          (panning on) an antibody-coated substrate such as a culture dish. CSA nucleic acids isolated by the method of the invention, and the proteins they encode, are useful for immunodiagnostic and immunotherapeutic applications, including the diagnosis and treatment of immune-mediated infections, diseases, and disorders in treatment of immune-mediated infections, diseases, and disorders in the contract of immune-mediated infections.
                                                                                                                                                                                                                                                                                          The present sequence is that of a human macrophage specific FCRI, as deduced from cDNA clone pl35 (see A50631), which was isolated from a cDNA library expressed in COS cells using a novel method of the invention designed to isolate CSA nucleic acids. The method is based upon transient expression of a CSA in eukaryotic cells and physical selection of cells expressing the antigen by adhesion to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25-FEB-1988;
13-JUL-1989;
23-MAR-1990;
13-JUL-1990;
Sequence
                                                                                                                                                                             animals, including humans.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Isolated nucleic acid molecule encoding the CD19 cell surface antigen, useful for immunodiagnosis and immunotherapy of immune-mediated infections or disorders, e.g. asthma, immune-complex disease, parasitic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 10; Column 53-55;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2000-586382/55
N-PSDB; A50631.
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.344 AA;
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88US-0160416.
89US-0379076.
90US-0498809.
90US-0553759.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note=
333
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220 ·
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268
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note=
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                                                                                                                                                                                                                                                                                                                                                                                                                                    75pp; English
                                                                                                                                                                         These disorders include asthma,
                                 es, is useful c ligand will n therapy.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CCT"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TTC"
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Query Match Matches

Local Similarity nes 11; Conser 1 EDSGLYQCEAAT 12

Conservative

89.1%; 91.7%;

Score 57; DB 2: Pred. No. 0.13; 0; Mismatches

DB 21; Length 344; 1; · Indels.

0;

0;

21-MAY-1992

(first entry)

0;

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R22550
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R20811
ID R3
XX
AC R2
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                                                                                                                                                                                                                                                                 Query Match
Best Local S
Matches 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This amino acid sequence was predicted from the cDNA sequence of cDNA clone p135. It differs from the sequence predicted from clones p98/X2 and p90 (see Q21179 and Q21180, respectively) at position 25; p135 encodes a Ser residue and the other two clones predict a Thr residue. At position 58, p135 predicts Leu and p90 predicts Val. Sequences predicted from all 3 clones show the typical features of a type I integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane-spanning domain, and a short, highly charged cytoplasmic domain. The extracellular portion contains six protein and allowed the contains and a short in the charged cytoplasmic domain. The extracellular portion contains six protein and a short in the charged cytoplasmic domain.
                                                                                                                                                                                                                                                                                                                                                                                                    potential N-linked glycosylation sites and six Cys residues distributed among three C2 set Ig-related domains. A fusion protein of FCRI and a receptor ligand will be helpful to increase the potency of antibodies in therapy.
                                                      R22550;
                                                                                        R22550 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Seed B,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 10; Page 94a; 160pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New CD53 cell surface antigen and DNA encoding immuno-therapy and diagnosis of haematopoietic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1992-056864/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO9201049-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rapid immunoselection cloning technique; cell surface antigen; immunodiagnosis; high affinity receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GEHO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human macrophage-specific FcRI receptor encoded by clone p135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21-MAY-1992 (first entry)
                                                                                                                                                                                           253
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                                                                                                                                                                                                                                                                   l Similarity 91.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                374
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                                                                                            Protein;
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                                                                                                                                                                                         264
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                                                                                                                                                                                                                                                                                    89.1%;
91.7%;
                                                                                          374 AA.
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                                                                                                                                                                                                                                                                                      Score 57; DB 13; Pred. No. 0.14;
                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           neoplasms, etc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     it
                                                                                                                                                                                                                                                                                                        Length 374;
                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                   0,
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RESULT
R91438
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Best Local Similarity 91.:
Matches 11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cDNA clone p30. It differs from the sequence predicted from clone p135 (see Q21178) at position 25; p135 encodes a Ser residue and clone p90 predicts a Thr residue. At position 58, p90 predicts val while clones p135 and p98/X2 (see Q21179) both predict Leu. Sequences predicted from all 3 clones show the typical features of a type I integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane-spanning domain, and a short, highly charged cytoplasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and six Cys residues distributed among three C2 set Ig-related domains. A fusion protein of FCRI and a receptor ligand will be believed the protein of FCRI and a receptor ligand will be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This amino acid sequence was predicted from the cDNA sequence of cDNA clone p90. It differs from the sequence predicted from clone p135 (see Q21178) at position 25; p135 encodes a Ser residucione p135 (see Q21178) at position 25; p135 encodes a Ser residucione p135 (see Q21178) at position 25; p135 encodes a Ser residucione p135 (see Q21178) at position 25; p135 encodes a Ser residucione p135 (see Q21178) at position 25; p135 encodes a Ser residucione p135 (see Q21178) at position 25; p135 encodes a Ser residucione p135 (see Q21178) at position 25; p135 encodes a Ser residucione p135 (see Q21178) at p135 encodes a Ser residucione p135 (see Q21178) at p135 encodes a Ser residucione p135 (see Q21178) at p135 encodes a Ser residucione p135 (see Q21178) at p135 encodes a Ser residucione p135 (see Q21178) at p135 encodes a Ser residucione p135 (see Q21178) at p135 encodes a Ser residucione p135 (see Q21178) at p135 encodes a Ser residucione p135 (see Q21178) at p135 encodes a Ser residucione p135 (see Q21178) at p135 encodes a Ser residucione p135 (see Q21178) at p135 encodes a Ser residucione p135 (see Q21178) at p135 encodes a Ser residucione p135 (see Q21178) at p135 encodes a Ser residucione p135 (see Q21178) at p135 encodes a Ser residucione p135 (see Q21178) at p135 encodes at p135 (see Q21178) at p135 encodes at p135 (see Q21178) at p135 encodes at p135 (see Q21178) at p135 (see Q21178) at p135 (see Q21178) at p135 (see Q21178) at p135 (see Q21178) at p135 (see Q21178) at p135 (see Q21178) at p135 (see Q21178) at p135 (see Q21178) at p135 (see Q21178) at p135 (see Q21178) at p135 (see Q21178) at p135 (see Q21178) at p135 (see Q21178) at p135 (see Q21178) at p135 (see Q21178) at p135 (see Q21178) at p135 (see Q21178) at p135 (see Q21178) at p135 (see Q21178) at p135 (see Q21178) at p135 (see Q21178) at p135 (see Q21178) at p135 (see Q21178) at p135 (see Q21178) at p135 (see Q21178) at p135 (see Q21178) at p135 (see Q21178) at p135 (see Q21178) at p135 (see Q21178) at p135 (see Q21178) at p135 (see Q21178) at p
                                                                                                                                                                                                                                                                                           Cell surface antigen; cloning; immunoselection; therapy; diagnosis; vector; FCRI; Fc receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R91438;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 R91438 standard; Protein; 374 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Rapid immunoselection cloning technique; cell surface antigen; immunodlagnosis; high affinity receptor.
       Misc-difference
                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                      Human FcRI (cDNA clone pl35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-OCT-1996 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New CD53 cell surface antigen and DNA encoding it - for immuno-therapy and diagnosis of haematopoietic neoplasms,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          N-PSDB; Q21180
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                                                                                                                  Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-JUL-1990;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         253
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 EDSGLYQCEAAT 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             edsglywceaat 264
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            increase the potency of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   90US-0553759
                                                                                                               Location/Qualifiers
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                                      Amiot M;
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91.7%;
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Pred. No. 0.14;
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                                                                                                                                                                                                                                                                                                                              immunotherapy;
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Best Local Similarity 91.7
Matches 11; Conservative
01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
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25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
18-OCT-1993;
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                                                                                                                                                                                                                      09-APR-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                     Cell surface antigen; cloning; immunoselection; immunotherapy; therapy; diagnosis; vector; FcRI; Fc receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human FcRI (cDNA clone p90 product).
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N-PSDB; T14717.
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92US-0983647.
88US-0160416.
89US-0379076.
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900S-0553759
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91.78;
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Pred. No. 0.
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DB 17;

Length 374; Indels

0

Gaps

0,

Aruffo A,

Seed

В;

(GEHO) GEN

HOSPITAL CORP

13-JUL-1990; 18-OCT-1993;

90US-0553759. 93US-0139273.

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Best Local S
Matches 11
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                     Misc-difference
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                                                                                                                                                                                                     Misc-difference
                                                                                                                                                                                                                                                    Macrophage; FcRI; cell surface antigen; human; immunoselection; panning; immunodiagnosis; diagnosis; immunotherapy; gene therapy immune disorder; infection; asthma; immune-complex disease; amyloidosis; multiple sclerosis; parasitic disease; Fc receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The amino acid sequence (W00859) of human FCRI was detd. from a cDNA clone, p90 (T14719), obtd. from a cDNA library using an immunoselection cloning method. FCRI is a high affinity receptor for the Fc portion of IgG, normally located on cell surfaces of macrophages. Other CDNA clones (see also T14717-18) coded for variants (R91438-39) of the FCRI sequence. A fusion protein of FCRI and a receptor ligand will be helpful in increasing the potency of antibodies in therapy.
                                                                                                                                                                                                                                     Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                               Y96134 standard; Protein; 374 AA.
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253 edsglywceaat 264
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                  /note=
                                                              /note=
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:51
/note=
116
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.01-DEC-1992;
25-FEB-1988;
.13-JUL-1989;
.23-MAR-1990;
.13-JUL-1990;
                       Stamenkovic I,
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N-PSDB;
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Misc-difference 140
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2000-586382/55
)B; A50592.
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89US-0379076
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216
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268
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126
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Isolated nucleic acid molecule encoding the CD19 cell surface antigen, useful for immunodiagnosis and immunotherapy of immune-mediated infections or disorders, e.g. asthma, immune-complex disease, parasiti immune-complex disease, parasitic

Example 10; 75pp; English.

proteins they encode, are useful immunotherapeutic applications, i treatment of immune-mediated infe based upon transient expression of a CSA in eukaryotic cells and physical selection of cells expressing the antigen by adhesion to (panning on) an antibody-coated substrate such as a culture dish. The present sequence is that of a human macrophage specific FCRI, as deduced from cDNA clone p135 (see A50592), which was isolated from a cDNA library expressed in COS cells using a novel method of the invention designed to isolate CSA nucleic acids. The method is animals; including humans. CSA nucleic acids isolated by the method of the invention, and amyloidosis, These disorders infections, l for immunodiagnostic and including the diagnosis a fections, diseases, and di include asthma diseases or multiple disorders in is

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                                                                                                                                                                                                                                                                                                                                                                         The present sequence is the human high affinity Fc receptor, FcgammaRI. FcgammaRI is also known as CD64. Fc receptors play an important role in defending the body against infections. First, pathogens are opsonised by serum immunoglobulins. The resulting complex then binds to
                                                     cells expressing Fc receptors. FcgammaRI molecules are expressed by monocytes and macrophages, but expression can also be induced on neutrophils and eosinophils. Upon Fc receptor activation, immune effector pathways are activated, leading to immune response. The present sequence may be modified to produce recombinant versions. The recombinant Fc receptor consist only of the extracellular portion of the receptor and are not glycosylated i.e. they do not have transmembrane domains or signal peptides. The recombinant proteins may be used in immunoassays to determine the immune status of patients with chronic diseases of the immune system, e.g. AIDS, systemic lupus erythematosus (SLE), multiple
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sclerosis. FCRI is a high affinity receptor for the Fc portion of IgG, normally located on the cell surfaces of macrophages. The ability to interfere with such bonding, or to cause it to occur on surfaces other than macrophages, is useful in therapy. A fusion protein of FCRI and a receptor ligand will be helpful to increase the potencies of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-367968/32
N-PSDB; A27466.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 03-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; high affinity Fc receptor; FcgammaRI; immunoglobulin; infection; immune response; CD64; monocyte; macrophage; neutrophil; eosinophil; HIV; 196; immunosuppressive; antirheumatic; cytostatic; antiinflammatory; AIDS; SLE; multiple myeloma; rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 26-28; 60pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel Fc receptor lacking transmembrane domains, a signal peptide, and glycosylation, useful for diagnosing and treating immune disorders and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (PLAC ) MAX PLANCK GES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      03-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EP1006183-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human high affinity Fc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11-SEP-2000 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EDSGLYQCEAAT 12
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(MM), or rheumatoid arthritis. In addition, pharmaceutical tions containing recombinant proteins may be used to treat o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  98EP-0122969.
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91.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    receptor, FcgammaRI.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              tumour.
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treat or
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Best Local Similarity
            antithyroid; antiallergic; antibacterial; antiviral; dermatological; neuroprotective; cardiant; thrombolytic; coagulant; nootropic; vasotropic; antipsoriatic and antiangiogenic. The polynucleotides and polypeptides can be used for preventing, treating or ameliorating medical conditions and diagnosing pathological conditions. Polynucleotides, polypeptides, antibodies, agonists and antagonists from the present invention may be used to treat immune disorders by activating or inhibiting the proliferation, differentiation or mobilisation of immune cells, to treat disorders of haematopoietic cells, autoimmune disorders, allergic reactions, graft versus host disease and organ rejection.
                                                                                                                                                                          B43398 to B44239. The proteins can have activities based on the tissues and cells the genes are expressed in Example of activities include: cytostatic; proliferative; vulnerary; immunomodulator; antidiabetic; antiasthmatic; antirheumatic; antiarthritic; antiinflammatory; antithyroid; antiallergic: antiharterial antimicallergic: antiharterial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          vasotropic; antipsoriatic; antiangiogenic; gene therapy; inflammation;
immune disorder; haematopoietic cell disorder; autoimmune disorder;
allergic reaction; graft versus host disease; organ rejection;
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allergic reactions,
modulate haemostatic
                                                                                                                                                                                                                                                                                                     Claim 11;
                                                                                                                                                                                                                                                                                                                                  Novel isolated nucleic acids comprising sequences encoding peptides useful for treating or diagnosing e.g. cancer - \,
                                                                                                                                                                                                                                                                                                                                                                                         N-PSDB; C77892.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-MAR-2000; 2000WO-US05882
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antidiabetic; antiasthmatic; antirheumatic; antiarthritic; antiviral;
antiinflammatory; antithyroid; antiallergic; antibacterial; cardiant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      diagnosis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human cancer associated protein sequence SEQ ID NO:1128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             08-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                B43683 standard; Protein; 399
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Rosen CA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             haemostatic; thrombolytic; cardiovascular disorder; infection;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cancer associated gene; cancer antigen; detection;
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                                                                                                                                                                                                                                                                                                     Page 1739-1740;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cytostatic; proliferative; vulnerary; immunomodulator;
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91.78;
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Pred. No.
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Best Local
01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
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                                                    21-MAY-1997;
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11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  399 AA;
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        92US-0983647.
88US-0160416.
89US-0379076.
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23
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antigens from mammalian lymphocytes (see W80440-55). The isolated genes can be expressed in a prokaryotic or eukaryotic host cells produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V63444) which allow the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and
                                                                                                                                                                                                             adhesion to an antibody-coated substrate. The method is useful for the isolation and cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V63442-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially human lymphocyte antigenic sequences
Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                 This is the amino acid sequence of human Fc receptor I (FCRI), as deduced from cDNA clone p135 (see V63456) isolated using a rapid immunoselection cloning method from a CDNA library expressed in COS cells. The cDNA library was constructed from polyA RNA of cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 10; Column 53-54; 75pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-MAY-1997;
                                                                                                                                                                                                                                                                                                                                            expression libraries is based on transient expression of an antigen
                                                                                                                                                                                                                                                                                                                                                                  were also obtained.
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                                                              treatment of immune-mediated
                                                                                                                                                                                                                                                                                                                                                                  from a single patient undergoing extracorporeal interleukin-2 induction therapy. Clones p90 (see X07372) and p98 (see X07373) were also obtained. A novel method for cloning cDNAs from mamma
                                                                                                                                                                                                                                                                                                                          in eukaryotic cells and selection of cells expressing
                                                                                  mmunotherapeutic applications,
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DB; V63456.
                                          including
  374 AA;
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                                              humans
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                                                                  disorders
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RESULT
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Best Local Similarity
Matches 10; Conserv
              Misc-difference
                                                                                  cloning
                                                                                        Fc receptor
                                                                                                        Human Fc receptor I
                                                                                                                       07-JUN-1999
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                             Misc-difference
                                            Misc-difference
                                                                  Homo sapiens
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                                                                                                                                                                                                                      Conservative
                                                                                                                      (first entry)
                                                                                         FCRI;
/note= "encoded by GAC" 45
               44
                              23
                                              Location/Qualifiers
                     'note= "encoded by CTC"
                                     'note=
                                                                                                                                                                                                       12
                                                                                                                                                                                                                              75.0%;
                                                                                         cell surface antigen; lymphocyte; human;
                                     "encoded
                                                                                                                                                                                                                      0;
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Pred. No. 3.8;
                                     ьу
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                                    'TGG"
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Misc-difference

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01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
21-MAY-1997;
                                                       adhesion to an antibody-coated substrate. The method is useful for the isolation and cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V63442-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated genes can be expressed in a prokaryotic or eukaryotic host cells to produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V6344) which allow the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and
                                                                                                                                                                                                                                                expression libraries is based on transient expression of an antigen in eukaryotic cells and selection of cells expressing the antigen by adhesion to an antibody-coated substrate. The method is useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New cloning vector and polylinker - based on efficient cloning and expression of mammalian
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N-PSDB; X07372.
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                                                                                                                                                                                                                                                                                                       This is the amino acid sequence of human Fc receptor. I (FCRI), as deduced from CDNA clone p90 (see X07372) isolated using a rapid immunoselection cloning method from a CDNA library expressed in COS cells. The cDNA library was constructed from polyA RNA of cells from a single patient undergoing extracorporeal interleukin-2 induction therapy. Clones p135 (see V63456) and p98 (see X07373) were also obtained. A novel method for cloning cDNAs from mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 10; Column 53-54; 75pp; English
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                     treatment of immune-mediated infections,
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                                        mmunotherapeutic applications,
including humans
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89US-0379076.
90US-0553759.
97US-0861205.
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                       diseases,
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                       and disorders of
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Sequence

374 AA

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Query Match
Best Local Similarity
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WPI; 1998-609251/51
N-PSDB; X07373.
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                           Aruffo A, Seed B;
                                                                                                                      21-may-1997;
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                                                               21-MAY-1997;
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89US-0379076.
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New cloning vector and polylinker – based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially human lymphocyte antigenic sequences

Example 10; Column 53-54; 75pp; English.

cc deduced from cDNA clone p98/X2 (see X07373) isolated using a rapid cc immunoselection cloning method from a cDNA library expressed in COS cells. The cDNA library was constructed from polya RNA of cells crom a single patient undergoing extracorporeal interleukin-2 from a single patient undergoing extracorporeal interleukin-2 clones also obtained. A novel method for cloning cDNAs from mammalian cc expression libraries is based on transient expression of an antipen by cathesion to an antibody-coated substrate. The method is useful for cd adhesion to an antibody-coated substrate. The method is useful for cd the isolation and cloning of any protein which can be expressed and cc transported to the cell surface membrane of a eukaryotic cell. It can be expressed to clone genes (see V63442-63) encoding cell surface can tigens from mammalian lymphocytes (see W80440-55). The isolated cc genes can be expressed in a prokaryotic or eukaryotic host cells to croduce the encoded protein. The invention also provides high cc efficiency expression vectors (see V63441 and V6344) which allow cc the generation of very large mammalian expression libraries. The cc purified genes and proteins are useful for immunodiagnostic and communotherapeutic applications, including the diagnosis and creatment of immune-mediated infections, diseases, and disorders of animals. Including high mamans animals, including humans.

Sequence 374 AA;

Query Match 75.1 Best Local Similarity 83.1 Matches 10; Conservative 75.0%; 83.3%; Score 48; DB 19; Length 374; Pred. No. 3.8; 0; Mismatches 2; Indels 0; Gaps

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Search completed: June 4, 2001, 12:13:31 Job time: 204 sec

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Post-processing: Minimum Match 0%
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Copyright (c) 1993 - 2000 Compugen Ltd.
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US-08-408-03-6
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US-08-714-901-6
US-08-928-383B-11
US-08-282-951-6
US-08-643-839-2
US-08-643-839-2
US-08-643-839-2
US-08-643-839-2
US-08-98-707A-2
PCT-US93-00031-11
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ALIGNMENTS

MOLECULE TYPE: US-08-752-307B-8 JS-08-752-307B-8 Patent No. INFORMATION FOR SEQ ID:NO:
SEQUENCE CHARACTERISTICS
LENGTH: 605 amino acid CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 19-NOV-19 ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette ATTORNEY/AGENT INFORMATION:
NAME: Meiklejohn, Ph.D., An
REGISTRATION NUMBER: 35,283
REFERENCE/DOCKET NUMBER: 09 CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: NUMBER OF SEQUENCES: TELECOMMUNICATION INFORMATION: TITLE OF INVENTION: APPLICANT: APPLICANT: APPLICANT: OPERATING SYSTEM: Windows95 SOFTWARE: FastSEQ for Windows Version TOPOLOGY: TELEFAX: FILING DATE: COMPUTER: COUNTRY: STREET: TELEPHONE: ADDRESSEE: . 8, Fr. No. 5952171 TMATT INFORMATION: 02110-2804 amino acid Application US/08752307B MA 605 amino acids 617-542-8906 225 Franklin Street SD McCarthy, Sean Gearing, David Levinson, IBM Compatible Fish & protein 19-NOV-1996 , Douglas A. METHOD FOR IDENTIFYING G Richardson, P.C. US/08/752,307B œ 09404/020001 Anita L. IDENTIFYING GENES 2.0 OR MEMBRANE-ASSOCIATED PROTEIN

Query Match Best Local S Matches 8

Similarity 8; Conserv

Conservative

75.0%;

Score 48: DB Pred. No. 2.1; 1; Mismatches

DB 2;

Length 605; Indels

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303 EDEGLYECEA 312

EDSGLYQCEA 10

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INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 1018 amin
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: protein
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                                                                                                                                                         APPLICANT: Bruskin, Arthur APPLICANT: Jarosz, David E.
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                                                                                                                                          APPLICANT:
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                                  CORRESPONDENCE ADDRESS:
                                                                                                         APPLICANT:
                                                                                                                        APPLICANT:
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TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-8864/9741
                                                   NUMBER OF SEQUENCES:
                                                                     TITLE OF INVENTION
                                                                                       APPLICANT:
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REGISTRATION NUMBER: 30,742
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3: Banner, Birch, McKie & Beckett
1001 G Street, N.W.
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                                                                   Vogelstein, Bert
Zabrecky, James R.
VENTION: Antibodies Specific for DCC Gene Product
                                                                                                                        Johnson, Karen
Kinzler, Kenneth W.
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N: 435
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
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MEDIUM TYPE: Floppy disk
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NAME: Kagan, Sarah A.
REGISTRATION NUMBER: 32,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/026,100
FILING DATE: 13-SEP-1996
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                                                                                                                                                                                 SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
TELECOMMUNICATION INFORMATION:
                                                           FILING DATE: 13-SEP-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                        ITTLE OF INVENTION: CAR, A No. 6210921el Coxsackievirus and Adenovirus
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CLASSIFICATION:
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                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DDS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE:
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                                NAME: Mandragouras, Amy E. REGISTRATION NUMBER: 36,20
                                                                                                                                                                                                                                                                                                                                           CITY: Boston
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              REFERENCE/DOCKET NUMBER:
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28 State Street
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Pred. No. 7
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TELEFAX: (617)7
NFORMATION FOR SEQ

(617)742-4214

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617)227-7400 ID NO

TELEPHONE:

SEQUENCE CHARACTERISTICS:

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 Query Match
Best Local Similarity
Matches 8; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  equence 94,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       tent No.
                                                                                                                                  VECTOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE:
FRAGMENT TYPE:
                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/896,457
FILING DATE: 27-MAY-1992
                                                                                                                                                                                                                                                      TTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                        OMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PPLICANT:
                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/O
FILING DATE: 31-OCT-1994
                                                                                            STRANDEDNESS:
                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 54:
                                                                                                                                                                                                                             NAME: BENT, Stephen A REGISTRATION NUMBER: :
                                                                                                                                                                                                                                                                                                                                                                                                                                                              STATE: D.C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      48
                                                                                                                                                                                                                                                                                                               CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY:
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                                                                                                                       LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION:
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                                                                                                       amino acids
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                                                                                                                                                               90413
                                                                                                                                                                                                                                             BENT,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3000 K Street, N.W., Suite 500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            75 amino acids
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                                                                             ss: single
                                                                                                                                                                                                                                                                                                                                                                 PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HULETT, Mark D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BONADONNA, Lisa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Foley & Lardner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       peptide
internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       71.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bernadette M
              71.98;
                                                                                                                                                                                                                                                                                                                                      US/08/332,562A
                                                                                                                                                94:
                                                                                                                                                                                                                 54270/119/GRHA
           Score 46; D
Pred. No. 0.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 4;
                         DB 2; Length 91;
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Indels
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Gaps
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Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                    US-08-752-307B-12
                                                                       Sequence 6,
                                          atent No. 5688916
GENERAL INFORMATION:
                                                                                       08-408-093-6
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NFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               tent No.
                                                                                                                                                                                                                                                                                                  TOPOLOGY: 1
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION: NAME: Meiklejohn, Ph.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: 435
PRIOR APPLICATION DATA:
 TITLE OF INVENTION:
                               APPLICANT:
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                                                                                                                                                 388 EDSGMYQCVA
                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: 617-542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: EILING DATE: 19-NO
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                                                                                                                                                                             1 EDSGLYQCEA 10
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5952171
                                                                                                                                                                                                                                                                                                                              amino acid
                                                                       Application US/08408093
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                                                                                                                                                                                                                                                                                                                                               607 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              McCarthy, Sean A. Gearing, David P.
            Hemperly,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Levinson,
                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                IBM Compatible
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-NOV-1996
                                                                                                                                                                                                                                                                                                protein
                               Robert A.
                                                                                                                                                                                                                         71.9%;
Human Cell Adhesion Molecule and Nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ph.D.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    METHOD FOR IDENTIFYING GENES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US/08/752,307B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   35,283
                                                                                                                                                                                                                                                                                                                                                                                                                                                      09404/020001
                                                                                                                                                                                                           1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Anita L.
                                                                                                                                                                                                                         Score 46;
Pred. No. 4
                                                                                                                                                                                                                                       DB 2;
                                                                                                                                                                                                                                     Length 607;
                                                                                                                                                                                                           0; Gaps
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Acid Sequences

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NAME: Fugit, Donna R.
REGISTRATION NUMBER: 32,135
REFERENCE/DOCKET NUMBER: P-:
NFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                      NAME/KEY: Domain
LOCATION: 604..657
OTHER INFORMATION:
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                              PEATURE:
NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RIOR APPLICATION DATA:
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RIGINAL SOURCE:
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                                                                                                      LOCATION: 809..857
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                            NAME/KEY: Domain LOCATION: 707..760 OTHER INFORMATION:
                                  OTHER
                                                LOCATION:
                                                              NAME/KEY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/408,093 FILING DATE: 21-MAR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: UZIP: 07417
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET:
                                                                                                                                                     NAME/KEY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Franklin Lakes
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                               KEY: Domain FION: 905..952 RINFORMATION:
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SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                    Domain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disulfide-bond 45..94
                                                                                                                                                                                                                                                                                                                                                               Disulfide-bond 506..563
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243..290
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138..191
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Floppy disk
/label= FLR
/note= "conserved core of fibronectin type
III-like repeat"
                                                                                       /label= FLR
/note= "conserved core of fibronectin type
III-like repeat" <
                                                                                                                                                                                /label= FLR
/note= "conserved core of fibronectin type
III-like repeat"
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/note= "conserved core of fibronectin type
III-like repeat"
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Best Local Similarity
Matches 7; Conserv
1 EDSGLYQCEA 10
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303 EDEGIYECEA 312
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OTHER INFORMATION:
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                                                                                                                                                                                                                                                                                         571
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                                                                         71.98;
70.08;
                                                                                                                                               //label= ASN-glycos
/note= "potential site of ASN-linked
glycosylation"
                                                                                                                                                                                                                                      /label= ASN-glycos
/note= "potential site of ASN-linked
glycosylation"
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/note= "potential site of ASN-linked
glycosylation"
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/note= "potential site of ASN-linked
glycosylation"
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/note= "potential site
glycosylation"
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/note= "potential site of ASN-linked
glycosylation"
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/note= "potential site of ASN-linked
glycosylation"
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/note= "potential site
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/note= "potential site of ASN-linked
                                                                       Score 46; DB 1;
Pred. No. 7.6;
                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              of ASN-linked
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                                                                                       Length 1018;
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RESULT 8 US-08-408-420A-6

Sequence 6, Application US/08408420A Patent No. 5731154
GENERAL INFORMATION:

APPLICANT:

Reid, Robert A. Hemperly, John J.

Human Cell Adhesion Molecule and Nucleic

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PEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Fugit, Donna R.
REGISTRATION NUMBER: 32,135
                                                                                                                                                                                                                                     LOCATION: 604..657
OTHER INFORMATION:
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OTHER INFORMATION:
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CLASSIFICATION:
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416..464
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332..371
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III-like repeat"
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/note= "conserved core of fibronectin type
III-like repeat"
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/note= "conserved core of fibronectin type
III-like repeat"
/label= FLR
/note= "conserved core of fibronectin type
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US-08-714-901-6

Patent No. 5739

Application US/08714901

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Query Match
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/note= "potential site
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/note= "potential site of ASN-linked
glycosylation"
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/note= "potential site of ASN-linked
                                                     Score 46; DB Pred. No. 7.6; 2; Mismatches
                                                                  DB 1;
7.6;
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                                                                                Length 1018;
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                                                        Indels
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REFERENCE/DOCKET NUMBER: NFORMATION FOR SEQ ID NO: 6:
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APPLICATION NUMBER: US/08/040,741
FILING DATE: 26 MAR 1993
ATTORNEY/AGENT INFORMATION:
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LOCATION:
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                                                                                                       NAME/KEY: Domain LOCATION: 707..760 OTHER INFORMATION:
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LOCATION: 809..857
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FILING DATE: 17-SEP-1996
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/note= "conserved core of fibronectin type
III-like repeat"
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III-like repeat"
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                                                                 71.9%;
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/note= "potential
glycosylation"
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/note= "conserved core of fibronectin type
III-like repeat"
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/note= "potential site of ASN-linked
glycosylation"
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/note= "potential site
glycosylation"
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/note= "potential
glycosylation"
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/note= "potential site of ASN-linked
glycosylation"
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/note= "potential site of ASN-linked
glycosylation"
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/note= "potential site of ASN-linked
glycosylation"
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/note= "potential site of ASN-linked
glycosylation"
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/note= "potential site of ASN-linked
glycosylation"
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                                                                Score 46;
Pred. No.
                                                      Mismatches
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7.6;
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US-08-040÷741-6
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                                                                                                                                                    LOCATION: 604..657
OTHER INFORMATION:
OTHER INFORMATION:
OTHER INFORMATION:
                                                         NAME/KEY: Domain
LOCATION: 707..760
OTHER INFORMATION:
OTHER INFORMATION:
OTHER INFORMATION:
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MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                                             EATURE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                         EATURE
              NAME/KEY:
LOCATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MBER OF SEQUENCES:
NAME/KEY: Domain LOCATION: 809..857 OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                        NAME/KEY:
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ADDRESSEE: and Company
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243..290
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138..191
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/note= "conserved core
III-like repeat".
                                                         /label= FLR
/note= "conserved core of fibronectin type
III-like repeat"
 /label= FLR
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OTHER INFORMATION:
US-08-040-741-6
                             Matches
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Best Local
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                             Similarity 7; Conserv
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                                         71.9%;
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/note= "potential site
glycosylation"
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/note= "potential site of ASN-linked
glycosylation"
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/note= "potential site of ASN-linked
glycosylation"
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/note= "potential site of ASN-linked
glycosylation"
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glycosylation"
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/note= "potential site of ASN-linked
glycosylation"
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/note= "potential site of ASN-linked
glycosylation"
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/note= "potential site of ASN-linked
glycosylation"
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/note= "potential site of ASN-linked
glycosylation"
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III-like repeat"
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III-like repeat"
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                                           Score 46;
Pred. No. 7
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RESULT 11 US-08-928-383B-11

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RESULT 12
US-08-282-951-6
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Patent No. 621092
                                                                                                                                                                                                                                         Sequence 6, Application US/08282951
                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
Matches 6; Conser
                                                                                                                                                                                                                         Patent No.
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NFORMATION FOR SEQ ID NO:
                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: Morrison & Foerster
                                                                                                                                    TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: DF
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                          APPLICANT:
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LENGTH: 72 amino acids
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APPLICANT: and Marshall S. Horwitz
BITLE OF INVENTION: CAR, A NO. 6210921el Coxsackievirus and Adenovirus
FITLE OF INVENTION: Receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 13-SEP-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                      NUMBER OF SEQUENCES:
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CLASSIFICATION:
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OPERATING SYSTEM: PC-DOS/MS-DOS
                                  STATE:
                                                                STREET:
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                                                                                ADDRESSEE:
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                                                                755 Page Mill Road
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(617)742-4214
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internal
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                                                                                                                                    PROTEINS
                                                                                                                                                        CLONING GENES: WHICH ENCODE CELL-SURFACE AND SECRETED
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Pred: No. 0
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Matches 6; Conserv
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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REFERENCE/DOCKET NUMBER: 37.00
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 813-5600
                                                                                 APPLICATION NUMBER: US 08/643,839 FILING DATE: 07-MAY-1996 ATTORNEY/AGENT INFORMATION:
                                                                                                                                         CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
             NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE,DOCKET NUMBER: A-63291-1/WHD
FELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                          UMBER OF SEQUENCES:
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86 EDAGIYRCQA 95
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/282,951 FILING DATE: 29-JUL-1994
                                                                                                                                                                               APPLICATION NUMBER: FILING DATE: HEREW
                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                  SOFTWARE: PatentIn Release #1.0, Version #1:30
                                                                                                                                                                                                                                                                                                                                            COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Flehr, Hohbach, Test, Albritton & Herbert STREET: Four, Embarcadero Center, Suite 3400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                  San Francisco
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                                                                                                                                                                                                                                                                                                                                                                 California
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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SYSTEM: PC-DOS/MS-DOS
(415)
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Pred. No.
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STRANDEDNESS:
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US-08-643-839-2
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NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 767 amino acids
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                                                                                                                                                                                                                                                                           NAME: Dreger, Walter H.

REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-63291/WHD
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
TELEFAX: (415) 398-3249
                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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COUNTRY: United St
                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/08/643,839 FILING DATE: 07-MAY-1996
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CITY: San Francisco
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EDSGLYQCEAAT 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                E: Flehr, Hohbach, Test, Albritton & Herbert Four Embarcadero Center, Suite 3400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   767
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                                                   Conservative
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58.3%;
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58.3%;
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                                               Score 43; DB Pred. No. 17; 2; Mismatches
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Pred. No.
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                                                                                 Length 767;
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Search completed: June Job time: 224 sec
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                                                                                                                                                                                                                                                                                                                       TELEPHONE: (202) 833-7533
NFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION: TELEPHONE: (202) 833-7533
                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 07/649,565
FILING DATE: 01-FEB-1991
APPLICATION NUMBER: U.S. 07/487,038
FILING DATE: 02-MAR-1990
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
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                                                                                   661 EDAGVYECES 670
                                                                                                                                                                                                                                                                                                                                                                                            REGISTRATION NUMBER: 33, REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: ASCII
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STREET: Suite 300
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                                                                                                                 1 EDSGLYQCEA 10
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United States of America
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                                                                                                                                                                                                                                                                                                          828 amino acids
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                                                                                                                                                     Conservative
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Pred. No. 19;
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Compugen Ltd.

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Post-processing: Minimum Match 0%
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Maximum DB
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   Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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	65.6	65.6	65.6	65.6	65.6	65.6	65.6		65.6	65.6	67.2	67.2	67.2	67.2	67.2
	1052 1173	1033	822	821	820	790	739	647	538	475	26926	1356	1015	518	500
	22	2	N	1	ν	N	N	N	N	N	μ	N	N	N	2
ALIG	B49120 T25893	S19247	B54846	TVMSBK	S17295	A39627	A41288	B41288	JC2457	A54879	I38344	JC1402	T32186	JC4024	S59795
ALIGNMENT															
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5	protein-tyrosine k hypothetical prote	cell adhesion prot	fibroblast growth	fibroblast growth	fibroblast growth	protein-tyrosine k	vascular cell adhe	vascular cell adhe	vascular cell adhe	pregnancy-specific	titin, cardiac mus	protein-tyrosine k	hypothetical prote	poliovirus recepto	hypothetical prote

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Fc gamma (IgG) receptor I-A (high affinity) precursor - human
N;Alternate names: CD64
C;Specles: Homo sapiens (man)
C;Date: 30-Dec:1991 #sequence_revision 06-Sep-1996 #text_change 21-Jul-2000
C;Accession: A39878; I70304; B41357; S03018; I57525
R;van.de Winkel, J.G.J.; Ernst, L.K.; Anderson, C.L.; Chiu, I.M.
J. Biol. Chem. 266, 13449-13455, 1991
A;Title: Gene organization of the human high affinity receptor for IgG, FcgammaRI (CD A;Reference number: A39878; MUID:91302383
A;Accession: A39878
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Cross-references: EMBL:X14355; NID:g31333; PIDN:CAA32536.1; PID:g31334 A;Note: the authors translated the codon ACT for residue 25cas Ala C;Superfamily: Fc gamma receptor 1; immunoglobulin homology C;Keywords: alternative splicing; immunoglobulin receptor; transmembrane protein F;117-170/Domain: immunoglobulin homology <IMM>
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A41357
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Molecule type: mRNA
A; Residues: 1-344 < ALL1>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             R;Allen, J.M.; Seed, B.
Science 243, 378-381, 1989
Science 243, 378-381, 1989
A;Title: Isolation and expression of functional high-affinity Fc receptor complementa A;Reference number: A41357; MUID:89100284
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Species: Homo sapiens (man)
C;Date: 03-Apr-1992 #sequence_revision 03-Apr-1992 #text_change 21-Jan-2000
C;Accession: A41357; S03019
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Molecule type: mRNA
A; Residues: 1-344 <ALL2>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         R;Allen, J.M.; Seed, B.
Nucleic Acids Res. 16, 11824, 1988
                                                                                                                                                                                                                                                                                            A39878
                                                                                                                                                                                                                                                                                                                   RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Accession: A41357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              N;Alternate names: CD64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     gamma (IgG) receptor I (high affinity)
                                                                                                                                                                                                                                                                                                                                                                                                    253 EDSGLYWCEAAT 264
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity 91. es 11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                       1 EDSGLYQCEAAT 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             89.18;
91.78;
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Pred. No. 0.016;
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 2;
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A; Molecule type: DNA A; Residues: 1-374 <VAN>

nces: EMBL:U55211; NID:g1305526; PID:g1305527; PIDN:AAA99159.1 neural cell adhesion molecule L1; fibronectin type III repeat homology

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A; Molecule type: mRNA
A; Residues: 1-24,'5', 26-337,'T', 339-374 <AL2>
A; Residues: 1-24,'5', 26-337,'T', 339-374 <AL2>
A; Residues: 1-24,'5', 26-337,'T', 339-374 <AL2>
A; Cross-references: EMBL:X14356; NID::g31331; PIDN:CAA32537.1;
A; Cross-references: Capacity of the company of the compa
                                                                                                                                                                                                                                                         C; Date: 11-Ja
C; Accession:
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                                                                                                                                                             submitted to the EMBL Data A; Description: Molecular cl
                                                                                                                                                                                                                             R; Giordano,
                                                                                                                                                                                                                                                                                                                                                        neural cell adhesion molecule L1 - N; Alternate names: E587 antigen
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                                                                A;Status:
                                                                                            A; Accession: T43027
                                                                                                                                 A; Reference number:
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Porges, A.J.; Redecha, P.B.; Doebele, R.;
Clin. Invest. 90, 2102-2109, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cross-references: GB:L03418; NID:g184840; PIDN:AAA36049.1; PID:g292169 Allen, J.M.; Seed, B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Accession: I70304
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Reference number: 157525; MUID:93204964
Accession: 157525
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cleic Acids Res. 16, 11824, 1988
Title: Nucleotide sequence of three cDNAs
Reference number: S03018; MUID:89098339
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   Residues:
                                                                                                                                                                                                                                                                                                                             Species: Carassius auratus (goldfish)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 117-170/Domain: immunoglobulin homology <IMM2>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Keywords: alternative splicing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Superfamily: Fc gamma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cross-references:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gene: GDB:FCGR1A; CD64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cross-references: GB:S57204; NID:g298692; PIDN:AAD13887.1; PID:g4261587
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Molecule type: DNA
Residues: 1-8 <RE2>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Status: translated from GB/EMBL/DDB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Molecule type: mRNA
Residues: 1-24,'S',
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ssidues: 1-24,'S',26-337,'T',339-374 <ALL>
coss-references: GB:X14356; GB:M21091; NID:g31331; PIDN:CAA32537.1; PID:g31332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               253
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                                                                                                                                                                                                                                                                                              11-Jan-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Two cis-DNA elements involved in myeloid-cell-specific expression
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                                                             preliminary; translated
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                                                                                                                                                                                                                             S.; Laessing,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11;
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î-1232 <GIO>
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                                                                                                                                                                                                                                                                                           #sequence_revision 11-Jan-2000 #text_change
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                                                                                                                                                             , U.; Lottspeich, F.; Stuermer, C.A.O. ta Library, April 1996 cloning of goldfish E587 antigen; a c
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ling site: carbohydrate
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0.018;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             <EXT>
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C;Superfamily: neura
C;Keywords: cell adl
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S05944
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                                                                                                                                        Matches
          303
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A;Cross-references: GB:M81846; NID:g212206; PIDN:AAA48924.1; A;Experimental source: embryo C;Superfamily: interleukin-1 receptor type I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gene 111, 239-243, 1992
A;Title: The chicken II-1 receptor: differential evolution
A:Reference number: JQ1526; MUID:92175529
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        interleukin-1 receptor I precursor - chicken
C;Species: Gallus gallus (chicken)
C;Date: 17-Aug-1992 #sequence_revision 17-Aug-1992 #text_change 29-Sep-1999
                                                                                                                                                          A; Rereach
A; Accession: Suzza
A; Molecule type: mRNA
A; Molecule type: mRNA
The fidnes: 1-1020 <GEN>
The fidnes: EMBL: X14943;
                                                                                                                                                                                                                                                                                                                                        neuronal cell surface protein F3 precursor - mouse
C;Species: Mus musculus (house mouse)
C;Date: 01-Dec-1989 #sequence_revision 01-Dec-1989 #text_change
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Keywords: cytokine receptor; transmembrane protein F;1-19/Domain: signal sequence #status-predicted <SIG> F;20-555/Product: interleukin-1 receptor I #status predicted <INT> F;339-359/Domain: transmembrane #status predicted <TRA>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R;Guida, S.; Heguy, A.; Melli,
Gene 111, 239-243, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C; Accession:
                                                                                      A; Map position: 15F
C;Superfamily: contactin; fibronectin type III repeat homology;
F;1-20/Domain: signal sequence #status predicted <SIG>
F;256-312/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                    C; Accession: S05944
R; Gennarini, G.; Ci
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Residues: 1-555 <GUI>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Molecule type: mRNA
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                                                                                                                                                                                                                                                 A; Reference number: S05944;
                                                                                                                                                                                                                                                                      A;Title: The mouse neuronal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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                 Query Match
Best Local
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nes 8; Conser
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8; Conserv
Similarity 8; Conserv
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                                                                                                                                                                                                                                                                                         775-788,
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88.9%;
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Score 48;
Pred. No.
1; Mismatc
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                                                                                                                                                                               NID: g50937; PIDN: CAA33075.1;
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Pred. No. 0.93
1; Mismatches
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Pred. No. 1.3;
2; Mismatches
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1.3;
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EDEGLYECEA EDSGLYQCEA 10

312

Conservative

Mismatches

Indels

0;

Gaps

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A; Title: The carbonic anhydrase domain of A; Reference number: A57112; MUID:95354206 A; Accession: A57112
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A;Residues: 1-1021 <F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R;Peles, E.; Nativ, M.; Campbell, P.L.; Sakurai, Cell 82, 251-260, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Status: preliminary; nucleic acid sequence not shown; not compared with conceptual tr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C; Date: 03-Nov-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ;Cross-references: EMBL:Y00813; NID:g63328; PIDN:CAA68753.1; Note: part of this sequence, including the amino end of the Bruemmendorf, T.; Wolff, J.M.; Frank, R.; Rathjen, F.G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ;Ranscht,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ;Alternate
                                                                                                                                                                                                                                                                    Superfamily: contactin; fibronectin type III repeat homology; immunoglobulin homology Keywords: blocked carboxyl end; glycoprotein; lipoprotein; phosphatidylinositol linka
                                                                                                                                                                                                                                                                                                                                                                     Residues: 1-970, 'S', 972-1000, 1090-1091, 'GVLAYSGF' <BRU>
Cross-references: GB:X14877; NID:g1708784; PIDN:CAA33018.1; PID:g63385
Note: the carboxy-end hydrophobic stretch is compatible with the consensus motif for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cell Biol. 107, 1561-1573, 1
                                                                                                                                                                                                                                                                                                             Comment: Fll is a chick neural cell surface-associated glycoprotein implicated in neu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local
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                                                                                                                                                                   983-1002/Domain: transmembrane #status predicted <TMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             uron 2, 1351-1361, 1989
Title: Neural cell recognition molecule F11; homology with fibronectin type III and Reference number: JU0094; MUID:90180453
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Species: Gallus gallus (chicken)
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      Matches
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Residues: 1-1091 <RAN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Reference number:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Accession: S01998
                                                                                                                                                003-1091/Domain:
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                                                                                                                                                                                                          982/Domain: extracellular #status predicted <EXT>
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                                                                                                                        091/Product: contactin #status predicted <MAT>
                         Local
                                                                                                                                                                                                                                                   Domain: signal
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                                                                                                                                                                                         /Domain: immunoglobulin
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      8,
                         Similarity
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8; Conser
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      Conservative
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                   75.0%;
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80.0%;
    1;
                                                                                                                                                                                         homology <IMM>
                     Score
Pred.
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Pred. No.
                                                                                                                                                                                                                                                 glycoprotein; lipoprotein;
tatus predicted <SIG>
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                   No.
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                         DB 2;
1.7;
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                                        Length 1091;
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  Indels
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                                                                                                       (Ser)
Gaps
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probable tumor suppressor - C. Species: Homo saplens (ma C. Date: 18-Oct-1991 #sequen C. Accession: A38442 R.Nigro, J.M.; Cho, K.R.; F. R.Nigro, J.M.; Cho, K.R.; F. R.
                           N;Alternate names: pE4 protein
C;Species: Rattus norvegicus (Norway rat)
C;Date: 02-Aug-1994 #sequence_revision 02
C;Accession: A54017; A61206
R;Chadeneau, C.; LeMoullac, B.; Denis, M.
 A; Title:
                                                                                                                                          RESULT
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                                                                                                                                                                                                                 YO.
                                                                                                                                                                                                                                                                                                                                 A; Residues: 1-267 <NIG>
A; Cross-references: GB:M63696; GB
C; Keywords: transmembrane protein
                                                                                                                                                                                                                                                                                                                                                                             A; Reference number: A38442;
A; Accession: A38442
A; Status: preliminary
A; Molecule type: DNA
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                  J. Biol.
                                                                                                            colon carcinoma-associated antigen pE4 precursor
                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Title: Scrambled exons
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A; Cross-references: EMBL: AF004840;
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                                                                                                                              A54017
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Status: preliminary; translated
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C; Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C; Date: 24-Mar-1999 #sequence_revision
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C; Species:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A38442
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Best Local
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A novel member of the immunoglobulin gene superfamily expressed in rat carci
                  Chem.
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9; Conser
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                  269,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          norvegicus (Norway rat)
                  15601-15605,
                                                                                                                                                                                                                                                                                                                                                 GB:M63696; GB:M63700;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10
                                                                                                                                                                                                                                                                    73.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   75.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Fearon,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Library,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          human (fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gao, M.; Feinleib,
                                                                                                                                                                                                                                                                                                                                                                                                                               MUID: 91121517
                               Denis, M.G
                  1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                    Score 47;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              E.R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 48;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NID: g3550957; PIDN: AAC34735.1; PID: g3550958
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             September 1998
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                                                            02-Aug-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24-Mar-1999 #text_change 05-Nov-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Kern,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NO
                                                                                                                                                                                                                                                                                                                                                 GB:M63702; GB:M63718;
                                                                                                                                                                                                                                                                       DB
0.7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SE
                                                            #text_change 01-Dec-1995
                                                                                                            rat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ruppert, J.M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 1256;
                                                                                                                                                                                                                                                                                   Length 267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                 GB:M63698
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A; Molecule type: mRNA
A; Residues: 1-1447 <HED>
                                                                                A;Title: The DCC gene product in cellular differentiation A;Reference number: A54100; MUID:95011532 A;Accession: A54100
                                                                                                                                            R; Hedrick, L.; Cho, K.R.; Fear
Genes Dev. 8, 1174-1183, 1994
                                                                                                                                                                                                                              N; Alternate names: colorectal C; Species: Homo sapiens (man)
                                                                                                                                                                                                                                                  tumor suppressor protein DCC precursor - human N; Alternate names: colorectal cancer suppressor
                                                                                                                                                                                                                                                                                                                                                                                    밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C; Keywords:
A;Cross-references: EMBL:X76132; NID:g453209; R;Fearon, E.R.; Cho, K.R.; Nigro, J.M.; Kern,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Superfamily: neural cell adhesion molecule L1; fibronectin type
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Molecule
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Note: C18F3.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Map position:
A; Introns: 34/3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ;Molecule type: protein
;Residues: 34-41, X',43-53 <CH2>
;Residues: 36-41, X',43-53 <CH2>
;Note: the residue at position 9 is suggested to be glycosylated
;Comment: This 66K protein is detected by monoclonal antibody E4
;Keywords: glycoprotein; membrane protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ;Experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ;Chadeneau, C.; Denis, M.G.; nt. J. Cancer 47, 903-908, 1; Title: Characterization, is:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ;Accession: A61206
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Title: Characterization, isolation and a Reference number: A61206; MUID:91184910
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Residues:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ;Residues: 1-1287 <DUZ>;Cross-references: EMBL:U50067; PIDN:AAA93439.2;Experimental source: strain Bristol N2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 bu,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cross-references: GB:L12025
                                                                                                                                                                                        Accession: A54100; A40098
                                                                                                                                                                                                           Date: 02-Aug-1994 #sequence_revision 02-Aug-1994 #text_change 05-Nov-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Molecule
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Reference number: Z20953
Accession: T30988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Description:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Species: Caenorhabditis elegans
Date: 22-Oct-1999 #sequence_revision 22-Oct-1999 #text_change 17-Mar-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 pothetical protein C18F3.2 - Caenorhabditis elegans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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                                                                                                                                                                                                                                                                                                                                                                                    168 DAGLYQCEA 176
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gattung, S.
to the EMBL Data Library, August 1999
tion: The sequence of C. elegans cosmid C18F3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     preliminary; translated from GB/EMBL/DDBJ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       T30988
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    IV
; 72/1; 131/2; 157/1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA
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58.3%;
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                                                                                                                                                                  Fearon, E.R.; Wu,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Blottiere,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 47; DB
Pred. No. 1.1;
3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 47;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            254/3; 303/1; 400/1; 495/1; 592/1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
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1.1;
                                                                                                                                                                      T.C.; Kinzler, K.W.; Vogelstein,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB
PIDN:CAA53735.1; PID:g453210 S.E.; Simons, J.W.; Ruppert, J.M.; Hamilto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     terminal sequencing of a rat colon carcin
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                                                                                                                                                                                                                                                        DCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 1287;
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                                                                                                                             colorectal tumorigenesis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     III repeat homology;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
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                       R; Sears, D.
J. Immunol.
                                                                                                                                                                    A; Title: Structure and mapping of the gene A; Reference number: A46480; MUID:92166399
                                                                                                                                                                                                                                     C;Accession: A46480; A43511
R;Osman, N.; Kozak, C.A.; M
                                                                                                                                                                                                                                                                          C;Species: Mus musculus (house mouse)
C;Date: 18-Jun-1993 #sequence_revision 18-Nov-1994 #text_change 21-Jan-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                В
    A; Title: Molecular
                                                                                      A; Residues: 1-404 <OSM>
                                                                                                                             A; Status: preliminary
                                                                                                                                                 A; Accession: A46480
                                                                                                                                                                                                                                                                                                                       N;Alternate
                                                                                                                                                                                                                                                                                                                                                              A46480
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                                                              A; Note: sequence extracted
                                                                                                          A; Molecule
                                                                                                                                                                                                                                   R;Osman, N.; Kozak,
                                                                                                                                                                                                                                                                                                                                           Fc gamma (IgG) receptor high affinity - mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Gene:
                                                                                                                                                                                                               Immunol. 148, 1570-1575,
                                                                                                                                                                                                                                                                                                                                                                                                                                                267
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D.W.; Osman, N

Tate,

from

and

expression of the mouse high affinity Fc receptor

for

NCBI backbone (NCBIN:85205, NCBIN:85208, B.; McKenzie, I.F.C.; Hogarth, P.M.

NCBIN:85212

144, 371-378, 1990

type: DNA

EDAGFYWCEVAT

278

names: high

affinity IgG receptor

McKenzie, I.F.; Hogarth,

ъ. Ж

encoding mouse

high affinity

FC

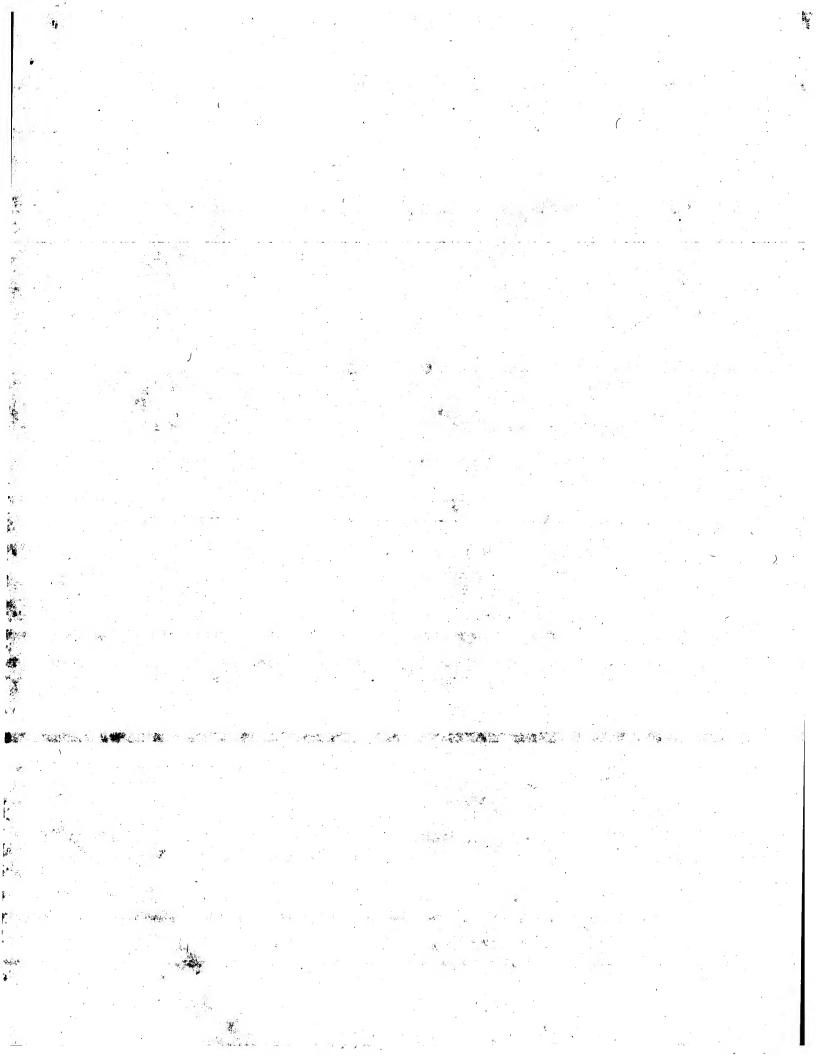
gamma

RI

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A; Molecule type: mRNA
A; Residues: 1-336 < RES>
A; Cross-references: EMBL:X70980; NID:g311748; PIDN:CAA50311.1;
C; Superfamily: Fc gamma receptor I; immunoglobulin homology
C; Keywords: immunoglobulin receptor
F;128-180/Domain: immunoglobulin homology < IMM>
                                                                                                                                                                                                                                                                     R;Prins, J.B.; Todd, J.A.; Rodrigues, N.R.; Gnosn, S.; Huyar Science 260, 695-698, 1993
A;Title: Linkage on chromosome 3 of autoimmune diabetes and A;Reference number: 148471; MUID:93242399
                                                                                                                                                                                                                                                                                                                                                 rc gamma (IgG) receptor high affinity - mouse
C;Species: Mus musculus (house mouse)
C;Date: 02-Jul-1996 #sequence_revision 02-Jul-1996 #text_change 21-Jan-2000
C;Accession: I48471
                                                                                                                                                                                                                                      A;Accession: I48471
A;Status: preliminary; translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             F;26-1447/Product: tumor suppressor protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C;Keywords: transmembrane protein; tumor suppressor F;1-25/Domain: signal sequence #status predicted <SIG>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Molecule type: mRNA
A; Residues: 1-750 <FEA>
A; Cross-references: GB:M32292; NID:g181492;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Science 247, 49-56, 1990 A; Title: Identification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Reference number: A40098; MUID:90100559
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Cross-references: GDB:119838;
A;Map position: 18q21.1-18q21.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C; Genetics:
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                                                      Query Match
Best Local
                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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1 EDSGLYQCEAAT
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                                   Similarity
8; Conserv
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8; Conservative
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                                   Conservative
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 12
                                                      71.98;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 47;
Pred. No.
                                                      Score 46;
Pred. No.
                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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3.4;
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                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               predicted
                                                                                                                                                                                                                                                                                                    defective Fc receptor for
                                                                                                                                                                                         PID:g311749
                                                                                                                                                                                                                                                                                                                                         P.M.;
                                       0;
                                       Gaps
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                                       0;
                                                                                                                                                                                                                                                                                                                                         L.S.; G
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A;Reference number: A43511; MUID:90111035 A;Accession: A43511

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Search completed: June 4, 2001, 12:15:44 Job time: 277 sec
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ₽
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   neural adhesion protein F3 precursor - bovine
C.Species: Bos primigenius taurus (cattle)
C.Date: 14-Nov-1995 #sequence_revision 08-Feb-1996 #text_change 20-Jun-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               R; Watanabe, K.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ene 160, 245-248, 1995
FILLE: Cloning of the cDNA encoding neural adhesion molecule F3 from bovine brain.
Reference number: JC4211; MUID:95369697
RACCESSION: JC4211
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ;Status: preliminary
;Molecule type: mRNA
;Residues: 1-404 <SEA>
                                                                                                                                                                                                                         Query Match 71.9
Best Local Similarity 70.0
Matches 7; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Experimental source: brain
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Residues: 1-1018 <WAT>
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303 EDEGIYECEA 312
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245-248, 1995
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Q06805	P55068	Q61361	P21802	P18460	095025	P79280	Q28343	Q61282	P07897	Q28062	P21804	
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L; X14356; C L; X14355; C S303018; S S303019; S A41357; B4 ; B41357; B4 P; B12319; I P; B12319; I 146760; - ; 146760; - ; 146760; - PF00047;	This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).	INITY RECEPTOR. CELLULAR LOCATION: TYPE I MEMBRANE PROTEI ERNATIVE PRODUCTS: 2 ISOFORMS; A' (SHOWN H DUCED BY ALTERNATIVE SPLICING. SUE SPECIFICITY: MONOCYTE/MACROPHAGE-SPEC ILARITY: CONTAINS 3 IMMUNOGLOBULIN-LIKE C ABASE: NAME=PROW; NOTE=CD guide CD64 entr -"http://www.ncbi.nlm.nih.gov/prow/cd/cd6		Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. NCBI_TaxID=9606; [1] SEQUENCE FROM N.A. MEDLINE=89098339; PubMed=2974947; Allen J.M., Seed B.; "Nucleotide sequence of three cDNAs for the human high affinity Fc	1 UMAN CG1_HUMAN STANDA 12314; P12315; 1-CCT-1989 (Rel. 12, 1-CCT-2000 (Rel. 40, 1-CCT-2000 (Rel. 40, 1-CCT-2000 (Rel. 40, 1-CCT-2000 (Rel. 40, CGR1A OR FCGR1 OR FC CGR1A OR FCGR1 OR FC CGR1A OR FCGR1 OR FC

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P14781; P10450;
01-MAR-1989 (Rel. 10, Created)
01-APR-1990 (Rel. 14, Last annotation
15-JUL-1999 (Rel. 38, Last annotation
                           MEDLINE=89286606;
Wolff J.M., Bruemn
                                                             GPI-ANCHOR
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MEDIINE-90180453; PubMed-2627374;

Bruemmendorf T., Wolff J.M., Rainer F., Rathjer F.G.;

"Neural cell recognition molecule F11: homology with fibronectin III and immunoglobulin type C domains.";

Neuron 2:1351-1361(1989).
                                                                                                                                                                        STRAIN=WHITE LEGHORN;
MEDLINE=89008597; Pub
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MEDLINE-89340657; PubMed=2474555;
Gennarini G., Cibelli G., Rougon G., Mattei M.-G., Goridis C.;
"The mouse neuronal cell surface protein F3: a phosphatidylinositol-anchored member of the immunoglobulin superfamily related to chicken contactin.";
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    J. Cell Biol. 109:775-788(1989).
    -I- FUNCTION: MEDIATES CELL SURFACE INTERACTIONS DURING NERVOUS

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P43146;
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01-NOV-1995
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                                                                                                                                              human esophageal squamous metastasis.";
                                                                                                                                                                                    Miyake S., Nagai K.,
                                                                                                                                                                                                                                                                       Cho K.R., Oliner J.D., Simons J.W., Hedrick L:, Preisinger A.C., Hedge P., Silverman G.A., Voge
                                                                                                                                                                                                                                                                                                                                             MEDLINE-91121517; PubMed-19
Nigro J.M., Cho K.R., Feard
Oliner J.D., Kinzler K.W.,
"Scrambled exons.";
                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE OF 107-472 FROM N.A. (SC MEDLINE-91121517; PubMed-1991322;
                                                                                                                                                                                                                                                                                                                                                                                                                                "Identification of a chromosome colorectal cancers.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE-90100559; PubMed-2294591; Fearon E.R., Cho K.R., Nigro J.M. Ruppert J.M., Hamilton S.R., Prei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A. MEDLINE-95011532;
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Mammalia; Eutheria;
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                                                                                                                                      Cancer Res.
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MEDLINE-94243823; PubMed-8187090;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE OF 1-750 FROM N.A.
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03 EDEGLYECEA 312
FUNCTION: IMPLICATED AS A TUMOR SUPPRESSOR GENE.
SUBCELLULAR LOCATION: TYPE I MEMBRANE PROTEIN
TISSUE SEBCIFICITY: FOUND IN AXONS OF THE CENTRAL AND PERIPP
NERVOUS SYSTEM AND IN DIFFERENTIATED CELL TYPES OF THE INTE:
DISBASE: COLORECTAL TUMORS THAT LOST THEIR CAPACITY TO
DIFFERENTIATE INTO MUCUS PRODUCING CELLS UNIFORMLY LACK DCC
EXPRESSION, INACTIVATION OF DCC DUE TO ALLELIC DELETION AND
POINT MUTATIONS MAY CAUSE BOTH LYMPHATIC AND HEMATOGENOUS
METASTASIS OF DESOPHAGEAL SQUAMOUS CELL CARCINOMAS.
SIMILARITY: CONTAINS 6 FIBRONECTIN TYPE III-LIKE DOMAINS.
                                                                                                                                                                         mutations
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Vogelstein B.;
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Pred. No.
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Preisinger
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tumor suppressor gene DCC
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Thomas G., Kinzler K.W.
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M32290; AAA52176.1;
M32290; AAA52177.1;
M63696; AAA52177.1;
M63700; AAA52179.1;
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SING (IN REF. 3).
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EXTRACELLULAR (POTENTIAL)
POTENTIAL.
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SIGNAL

Glycoprotein; Immunoglobulin Anti-oncogene; Alternative in

in domain; Transmembrane; Signal; initiation; Alternative splicing.

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PRINTS;

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InterPro; IPR003006; -
Ffam; PF00041; fn3; 6.
Pfam; PF00047; ig; 4
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HSSP; P56276; 1TLK.
MGD; MGI:94869; DCC.
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Submitted (JUN-1996) to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cooper H.M., Armes P., Britto J., ("Cloning of the mouse homologue of gene (mDCC) and its expression in Oncogene 11:2243-2254(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRAIN=BALB/C; TISSUE=Brain;
MEDLINE=96112625; PubMed=8570174;
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Mammalia; J
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           titles requires a license agreement (S send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TISSUE SPECIFICITY: IN THE EMBRYO, EXPRESSED AT HIGH LEVELS I DEVELOPING BRAIN AND NEURAL TUBE. IN ADULT, HIGHLY EXPRESSED BRAIN WITH VERY LOW LEVELS FOUND IN TESTIS, HEART AND THYMUS. DEVELOPMENTAL STRACE: LOW LEVELS IN EARLY GESTATION. HIGHEST I EXPRESSED DURING MID GESTATION. LEVELS DECREASE IN LATE GESTA AND REMAIN AT THIS LEVEL IN THE ADULT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              mitted (JUN-1996) to the EMBL/GenBank/DDBJ databases. FUNCTION: IMPLICATED AS A TUMOR SUPPRESSOR GENE. SUBCELLULAR LOCATION: TYPE I TUMOR FORMSOR FROTEIN. ARE PRODUCTS: TWO FORMS OF THE PROTEIN ARE PRODUCED THE SAME GENE BY THE USE OF ALTERNATIVE INITIATION SITES. A FORM WHICH IS EXPRESSED ONLY IN THE EMBRYO IS PRODUCED BY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SIMILARITY: CONTAINS 4 IMMUNOGLOBULIN-LIKE C2-TYPE DOMAINS: SIMILARITY: CONTAINS 6 FIBRONECTIN TYPE III-LIKE DOMAINS:
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in the developing mouse embryo.";
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                                                                                                                                                                                                                                                                                                                                                                                Mus musculus (Mouse;
Eukaryota; Metazoa;
Mammalia; Eutheria;
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HIGH AFFINITY IMMUNOGLOBULIN GAMMA FC
RI) (FCRI) (IGG FC RECEPTOR I)
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Sears D.W., Osman N., Tate B., McKenzie
"Molecular cloning and expression of the
receptor for IgG.";
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           European Bioinformatics Institute by non-profit institute.
                                                                     FUNCTION: BINDS TO THE FC REGION OF IMMUNOGLOBULINS AFFINITY RECEPTOR.
SUBCELLULAR LOCATION: TYPE I MEMBRANE PROTEIN.
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TISSUE SPECIFICITY: MACROPHAGE-SPECICIC.
CAUTION: IT IS UNCERTAIN WHETHER MET-1 OR MET-10 IS
                                         SWISS-PROT entry is copyright. It is produced through a collaboration sen the Swiss Institute of Bioinformatics and the EMBL outstation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DOMAIN
                                                                                                                            MEDIINE-95048335; PubMed-7959734; Berglund E.O., Ranscht B.; "Molecular cloning and in situ localization gene (CNTNL) on chromosome 12q11-q12."; Genomics 21:571-582(1994).
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                                                                       molecule contactin.
                                                                                Reid R.A., Hemperly J.J.; "Identification and characterization
                                                                                                                                                                                                             Eukaryota; Metazoa;
Mammalia; Eutheria;
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PIR; A43511; A43511.
                                                                                                           SEQUENCE FROM N.A.,
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IMMUNOGLOBULIN-LIKE C2-TYPE DOMAINS FIBRONECTIN TYPE III-LIKE DOMAINS.
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Pred: No. 0.22
1; Mismatches
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Catarrhini; Hominidae;
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Matches

Similarity 80.08; Conservative

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Mismatches

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Indels

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Gaps

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73.48; 80.08;

Score 47; DB 1 Pred. No. 0.51;

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Length 1447;

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SWISS-PROT entry is

ROT entry is copyright. It is produced through Swiss Institute of Bioinformatics and the EN Bioinformatics Institute. There are no restr

There are no restrictions ng as its content is in

EMBL outstation a collaboration

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EMBL; Z21488;
HSSP; P40189;
MIM; 600016; -
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Pfam; PF00047; ig; 6.
Immunoglobulin domain; Glycoprotein; Cell adhesion; Repeat; Alternative specific control of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state o
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InterPro; IPR003006;
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EMBL; U07820;
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IG-LIKE C2-TYPE DOMAIN.

raniata; Vertebrata; Euteleostomi;
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Pfam; PF00041; fn3; 3.
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Immunoglobulin domain; (Cell adhesion; Repeat.
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"The axonally secreted cell adhesion molecule, axonin-1. Primary
structure, immunoglobulin-like and fibronectin-type-III-like domains
and glycosyl-phosphatidylinositol anchorage.";
Eur. J. Biochem. 204:453-463(1992).
                                                                                                                                                                                                                                                                                                                                                                         DOMAIN
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SUBCELLULAR LOCATION: ATTACHED TO THE NEURONAL GPI-ANCHOR.
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Osterwalder T., Inan C
                                                                                                                                                                                                                                                           REMOVED IN MATURE FORM.

IG-LIKE C2-TYPE DOMAIN.

IG-LIKE C2-TYPE TOMAIN.

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Sonderegger P.
                                                           Signal; GPI-anchor;
                       (GLCNAC.
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V TYPE-III.
V TYPE-III.
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4 CRC64;
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HE AXCAM L1 (G
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MEDIJNE-941400354; PubMed-8307567;
TSiotra C.P., Karagogeos D., Theodorakis K., Michaelidis M. Modi W.S., Furley J.A., Jessel M.T., Papamatheakis J.;
Modi W.S., Furley J.A., Jessel M.T., Papamatheakis J.;
"Isolation of the cDNA and chromosomal localization of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TAX1 OR TAG1.

Homo sapiens (Human).

Homo sapiens (Human).

Horota; Metazoa; Chordata;

Horota; Primates;
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15-JUL-1999 (Rel. 3
AXONIN-1 PRECURSOR
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Pfam; PF00041; fn3;
Pfam; PF00047; 1g; 6
                                                                                                                                                                                                                                                                               EMBL; X68274; CAA48335.1;
EMBL; X67734; CAA47963.1;
                                                                                                                                                                                                                                                                                                                    use by non-profit institutions as long modified and this statement is not removed entitles requires a license agreement (See or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (TAX1) encoding the human axonal glycoprotein TAG-1. Genomics 18:562-567(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE-93145965; PubMed-8425542;
Hasler T.H., Rader C., Stoeckli E.T.,
"CDNA Cloning, structural features, ar
human TAG-1/axonin-1.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FUNCTION: MAY PLAY A ROLE IN THE INITIAL GROWTH AND GUIDANCE
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(AXONAL GLYCOPROTEIN TAG-1)
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FIBRONECTIN TYPE-III.
FIBRONECTIN TYPE-III.
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CARBOHYD
                            InterPro;
InterPro;
                                                                                                                                                                                                                              "The axonal glycoprotein TAG-1 is an immunoglobulin superfamily member with neurite outgrowth promoting activity."; Cell 61:157-170(1990).

Cell 61:157-170(1990).

-i- FUNCTION: MAY PLAY A ROLE IN THE INITIAL GROWTH AND GUIDANCE AXONS. MAY BE INVOLVED IN CELL ADHESION.

-i- SUBCELLULAR LOCATION: ATTACHED TO THE NEURONAL MEMBRANE BY A
                                                                                                                                                                                                                                                                                                                                                                                                               P22063;
01-AUG-1991 (Rel. 19, Created)
01-AUG-1991 (Rel. 19, Last seg
15-JUL-1999 (Rel. 38, Last ann
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         Pfam; PF00041; fn3; 4. Pfam; PF00047; 1g; 6.
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                                                EMBL; M31725; AAA42201.1; PIR; A34695; A34695.
                                                                              or send an email to license@isb-sib.ch).
                                                                                         entities requires a license agreement
                                                                                                                    the European Bioinformatics Institute.
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                                                                                                                                                                                                                                                                                                          MEDLINE=90199890; P
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Mammalia; Eutheria;
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                                                                                                                                                                     DEVELOPMENTAL STAGE: TRANSIENTLY EXPRESSED IN THE DEVELOPING RAT NERVOUS SYSTEM.
SIMILARITY: CONTAINS 6 IMMUNOGLOBULIN-LIKE
                                                                                                                                                                                                   GPI-ANCHOR AND IS ALSO RELEASED FROM NEURONS
TISSUE SPECIFICITY: IN NEURAL TISSUES IN EMB
BRAIN, SPINAL CORD AND CEREBELLUM.
                                                                                                                                                            SIMILARITY: CONTAINS 6 IMMUNOGLOBULIN-LIKE C2-TYPE DOMAINS SIMILARITY: CONTAINS 4 FIBRONECTIN TYPE III-LIKE DOMAINS.
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8; Conser
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                           IPR001777; -. IPR003006; -.
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Rodentia;
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GLYCOPROTEIN TAG-1)
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CELL ATTACHMENT SITE
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STRAIN-129; TISSUE-Embryo;
MEDLINE-94117008; PubMed-7507076;
Cybulsky M.I., Allan-Motamed M., (
"Structure of the murine VCAM1 ger
                                                                                                                                                           TISSUE=Lymph node;
MEDLINE=93246254; PubMed=7683304;
Araki M., Araki K., Vassalli P.;
"Cloning and sequencing of mouse of Gene 126:261-264(1993).
                                                                                                                                                                                                                                                                                                                                                    MEDLINE=92181437; PubMed=1371918;
Hession C., Moy P., Tizard R., Chisholm P.
Burkly L., Miyake K., Kincade P., Lobb R.;
"Cloning of murine and rat vascular cell a
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STRAIN=FVB; TISSUE=Lung;
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Eukaryota; Metazoa; Chordata;
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KOTENAGA R., ANGO J., TSUBOI H., KAMIYA A.;

KOTENAGA R., ANGO J., TSUBOI H., KAMIYA A.;

SUBMITTED RECOGNITION. APPEARS TO FUNCTION

IN LEUKCCYTE-ENDOTHELIAL CELL ADHESION. INTERACTS WITH THE BETA-1

INTEGRIN VLA4 ON LEUKCCYTES, AND MEDIATES BOTH ADHESION AND SIGNAL

TRANSDUCTION. THE VCAMI/VLA4 INTERACTION MAY PLAY A

PATHOPHYSIOLOGIC ROLE BOTH IN IMMUNE RESPONSES AND IN LEUKCCYTE

EMIGRATION TO SITES OF INFLAMMATION.

1- SUBCELLULAR LOCATION: TYPE I MEMBRANE PROTEIN (LONG ISOFORM) OR

ATTACHED TO THE MEMBRANE BY A GPI-ANCHOR (SHORT ISOFORM).

1- ALTERNATIVE PRODUCTS: THE GPI-ANCHORED AND THE TRANSMEMBRANE

1- ISOFORMS ARE PRODUCED BY ALTERNATIVE SPLICING.

1- TISSUE SPECIFICITY: EXPRESSED ON INFLAMED VASCULAR ENDOTHELIUM, AS
                                                                                                                                                                                                                                                                                                                                                                                                                       This SWISS-PROT entry is copyright. It is produced between the Swiss Institute of Bioinformatics and the European Bioinformatics Institute. There are n
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Rumar A. G., Dal A...,

Ballantyne C.M.;

"Murine VCAM-1. Molecular cloning,

"Murine VCAM-1.";
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J. Biol.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               -!- SIMILARITY: CONTAINS 7 IMMUNOGLOBULIN-LIKE C2-TYPE DOMAINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Terry R.W., Kwee L., Levine J.E., Lawon ...... "Cytokine induction of an alternatively spliced murine "Cytokine induction of an alternatively spliced murine "Cytokine induction of an alternatively spliced murine "Cytokine induction of an alternatively spliced murine "Cytokine induction of an alternatively spliced murine "Cytokine induction of an alternatively spliced murine "Cytokine induction of an alternatively spliced murine "Cytokine induction of an alternatively spliced murine "Cytokine induction of an alternatively spliced murine "Cytokine induction of an alternatively spliced murine "Cytokine induction of an alternatively spliced murine "Cytokine induction of an alternatively spliced murine" "Cytokine induction of an alternatively spliced murine "Cytokine induction" of an alternatively spliced murine "Cytokine induction" of an alternatively spliced murine "Cytokine induction" of an alternatively spliced murine "Cytokine" of a cytokine induction of an alternatively spliced murine "Cytokine" of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of a cytokine induction of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRAIN=FVB/N; TISSUE=Kidney; MEDLINE=93317595; PubMed=7687058;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=95015899;
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ning of an inflammation-specific phosphatidyl i
of murine vascular cell adhesion molecule-1.",
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            L AS ON MACROPHAGE-LIKE INFLAMED TISSUE.
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9; PubMed=7523515;
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EMBL; L12541; AAC37607.
EMBL; U42327; AAB88576.
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Pfam; PF00047; ig; 5.
Immunoglobulin domain; Glycoprotein; Cell adhesion; Transmembrane;
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PIR; JN0581; JN0581.
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661 QDAGIYECESKT 672
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U12876; AAA80014.
U12873; AAA80015.
U12877; AAA80016.
                                                                                                                         Similarity 50 | 6; Conservative
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25, Created)25, Last sequence update)39, Last annotation update)
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IG-LIKE C2-TYPE DOMAIN 5.

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IG-LIKE C2-TYPE DOMAIN 7.
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MISSING (IN GPI-ANCHORED ISOFORM)
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N-LINKED (GLCNAC. . .) (POTENTIAL).
N-LINKED (GLCNAC. . .) (POTENTIAL).
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N-LINKED (GLCNAC. . . .) (POTENTIAL).
EKPFIVDISPGSÖVAAQVGDSVVLTCAAIGCDSPSF
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                                                                                                                                                                                                                                                                                                                                                                                            REMOVED IN MATURE GPI-ANCHORED ISOFORM
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                                      PRT;
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RESULT 13

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Burkly L., Miyake K., Kincade P., Lobb R.;
Burkly L., Miyake K., Kincade P., Lobb R.;
Cloning of murine and rat vascular cell adhesion molecule-1.";
Biochem. Biophys. Res. Commun. 183:163-169(1992).
-i. FUNCTION: IMPORTANT IN CELL-CELL RECOGNITION. INTERACTS WITH THE BETA-1
INTEGRIN VLA4 ON LEUKOCYTES, AND MEDIATES BOTH ADHESION AND SIGNAL
TRANSDUCTION. THE VCAMI/VLA4 INTERACTION MAY PLAY A
PATHOPHYSIOLOGIC ROLE BOTH IN IMMUNE RESPONSES AND IN LEUKOCYTE
EMIGRATION TO SITES OF INFLAMMATION.
-i. SUBCELLULAR LOCATION: TYPE I MEMBRANE PROTEIN.
-i. SUBCELLULAR LOCATION: TYPE I MEMBRANE PROTEIN.
-i. TISSUE SPECIFICITY: EXPRESSED ON INFLAMED VASCULAR ENDOTHELIUM, AS
                                                                                             CARBOHYD
CARBOHYD
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                                                                                SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      use by non-profit institutions as long as its cont modified and this statement is not removed. Usage by entities requires a license agreement (See http://www.or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            VASCULAR CELL ADHESION PROTEIN 1 PRECURSOR (V-CAM VCAM1 OR VCAM-1.
                                                                                                                                                                CARBOHYD
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                                                                                                                                                                                                                                                                                                                           DOMAIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EMBL; M84488; AAA42332.1;
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h 68.8%;
Similarity 50.0%;
6; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       JS0675.
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                                                                                                                                                                                                                                                                                                                                                                                    domain; Glycoprotein; Cell adhesion; Transmembrane;
                                                                              198
3287
379
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575
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739
698
720
739
91
                                                                                 MW.
                                                                                                                                                                          CYTOPLASMIC (POTENTIAL).

IG-LIKE C2-TYPE DOMAIN 1.

IG-LIKE C2-TYPE DOMAIN 2.

IG-LIKE C2-TYPE DOMAIN 3.

IG-LIKE C2-TYPE DOMAIN 4.

IG-LIKE C2-TYPE DOMAIN 6.

IG-LIKE C2-TYPE DOMAIN 6.

IG-LIKE C2-TYPE DOMAIN 7.
   5
Score 44; DB
Pred. No. 0.96
5; Mismatches
                                                                                                            N-LINKED
N-LINKED
                                                                                                                        N-LINKED
                                                                                                                                                                                                                                                                                                                          VASCULAR CELL
EXTRACELLULAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Rattus
                                                                                 5C608E5A1A1B100C
                                                                                               (GLCNAC. .
                DB 1;
0.96;
                                                                                                                                                                                                                                                                                                                          (POTENTIAL).
                                                                                                                                                                                                                                                                                                                                            ADHESION PROTEIN 1
   1;
                              Length 739;
                                                                                CRC64;
   Indels
                                                                                             (POTENTIAL).
(POTENTIAL).
(POTENTIAL).
(POTENTIAL).
(POTENTIAL).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .isb-sib.ch/announce/
   0;
   Gaps
   0;
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035136; 035962; 15-JUL-1998 (Re

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DISULFID
DISULFID
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                                                                                                                                                                                                                    EMBL; AF001287; AAB69125.1; -.
EMBL; AF001286; AAB69124.1; -.
EMBL; AF016619; AAC53375.1; -.
MGD; MGI:1095738; Ocam.
                                                                                                                   DOMAIN
                                                                                                                           TRANSMEM
                                                                                                                                      DOMAIN
                                                                                                                                                       SIGNAL
                                                                                                                                                                           Cell adhesion;
                                                                                                                                                                                                                                                                                                   This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Enthe European Bioinformatics Institute. There are no restricted the European Bioinformatics Institute.
                                                                                                                                                                                          Pfam; PF00041; fn3;
                                                                                                                                                                                                   InterPro; IPR001777; -. InterPro; IPR003006; -.
                                                                                                                                                                                                                                                                      or send
                                                                                                                                                               Immunoglobulin
                                                                                                                                                                                                                                                                                entities
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mus musculus (Mouse)
Eukaryota; Metazoa; (
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRAIN-C57BL/6J; TISSUE-Olfactory epithelium; MEDLINE-97476194; PubMed-9334170;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A. (LONG AND SHORT FORMS).
STRAIN-BALB/C; TISSUE-Olfactory neuroepithelium;
MEDLINE-97368238; PubMed-9221781;
YOShihara Y., Kawasaki M., Tamada A., Fujita H.,
Kagamiyama H., Mori K.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               axons.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15-JUL-1998 (Rel. 36, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
15-JUL-1998 (Rel. 36, Last annotation update)
NEURAL CELL ADHESION MOLECULE 2 PRECURSOR (N-CAM 2)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                J. Neurosci. 17:5830-5842(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "OCAM: A new member of the neural cell adhesion related to zone-to-zone projection of olfactory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADHESION MOLECULE) (R4B NCAM2 OR OCAM OR RNCAM.
                                                                                                                                                                                                                                                                     an
                                                                                                                                                                                                                                                                          non-profit institutions as long and this statement is not removed requires a license agreement (See
 1
20
698
719
129
225
315
415
491
136
232
322
322
                                                                                                                                                                                                                                                                  email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 N.A.
                                                                                                                                                              domain;
                                                                                                                                                                        Transmembrane;
                                                                                                                                                                                  ig; 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Chordata;
Rodentia;
                                                                                                                                                             Signa:
                                                   CYTOPLASMIC (POTENTIAL).
IG-LIKE C2-TYPE DOMAIN.
FIBRONECTIN TYPE-III.
        PROBABLE
                  PROBABLE
                           PROBABLE.
                                   PROBABLE
                                              FIBRONECTIN
                                                                                                                              NEURAL CELL ADHESION MOLEC EXTRACELLULAR (POTENTIAL).
                                                                                                                         POTENTIAL
                                                                                                                                                     POTENTIAL.
                                                                                                                                                        Glycoprotein; Repeat;
; GPI-anchor; Alternative splicing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Craniata; Veri
Sciurognathi;
                                                                                                                                        CELL ADHESION MOLECULE 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Fujita H., Hayashi
                                                                                                                                                                                                                                                                                        There are no rest
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Vertebrata; |
thi; Muridae;
                                                                                                                                                                                                                                                                       http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                  Usage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              molecule
and vomer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ecule family vomeronasal
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Murinae; Mus.
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NCM2_HUMAN
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                             Genomics 43:43-51(1997)
Genomics 43:43-51(1997)
-1- FUNCTION: MAY PLAY IMPORTANT ROLES IN SELECTIVE PASCICULATION
ZONE-TO-ZONE PROJECTION OF THE PRIMARY OLFACTORY AXONS.
-1- SUBCELLULAR LOCATION: TYPE I MEMBRANE PROTEIN.
-1- SUBCELLULAR LOCATION: EXPRESSED MOST STRONGLY IN ADULT AND FETAL
                                                        Pfam; PF00041; fn3; 2. Pfam; PF00047; ig; 5:
      Immunoglobulin
                                           Cell adhesion;
                                                                                                                                                                                                                                                                  This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its
                                                                                                                                           MIM; 602040
                                                                                                                                                                EMBL;
                                                                                                                                                                                           entities requires a license agreement (or send an email to license@isb-sib.ch)
                                                                                                                                                                                                                                     use by non-profit institutions as long modified and this statement is not removed
                                                                                                      InterPro;
                                                                                                                         InterPro;
                                                                                                                                                                                                                                                                                                                                                                                   -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo:sapiens (Human).
Homo:sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCM2_HUI
015394;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Down
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    that maps to chromosome region 21q21 and Down syndrome.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Paoloni-Giacobino A., Chen H., Antonarakis S.E.;
"Cloning of a novel human neural cell adhesion molecule gene (NCAM2)
that maps to chromosome region 21q21 and is potentially involved in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-JUL-1998 (Rel. 36, Last sequence update)
15-JUL-1998 (Rel. 36, Last annotation update)
NEURAL CELL ADHESION MOLECULE 2 PRECURSOR (N-CAM
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CARBOHYD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=97369930;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "ISSUE=Brain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EQUENCE FROM N.A.
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SIMILARITY:
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                                                                                                                                                        U75330; AAB80803.1;
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8; Conserv
                                                                                               IPR001777; -. IPR003006; -.
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domain;
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              Transmembrane; Glycoprotein; Repeat; domain; Signal.
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2 FIBRONECTIN TYPE III-LIKE DOMAINS
POTENTIAL
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N-LINKED
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Pred. No.
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                                                                                                                                                                                                              (See http://www.isb-sib.ch/announce/
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P35968; O60723; O1-JUN-1994 (Rel
01-CCT-2000 (Rel
01-CCT-2000 (Rel
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_HUMAN
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CARBOHYD
SEQUENCE
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TRANSMEM
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DOMAIN
                Oncogene
                                                              TISSUE-Umbilical vein;
MEDLINE-92019839; PubMed-1656371;
                                                                                                         TISSUE-Umbilical
Yu Y., Whitney R
"Coding region fo
                                                                                                                                                                                                                                 01-JUN-1994 (Rel. 29, Created)
01-OCT-2000 (Rel. 40, Last sequence update)
01-OCT-2000 (Rel. 40, Last annotation update)
01-OCT-2000 (Rel. 40, Last annotation update)
VASCULAR ENDOTHELIAL GROWTH FACTOR RECEPTOR 2 PRECURSOR (EC (VEGFR-2) (KINASE INSERT DOMAIN RECEPTOR) (FIK-1).
                                                                                                                                                                                                                                                                                                                                                                                                                              CARBOHYD
                                                         Terman
                                                                                                                                                   Submitted
                                                                                                                                                                                                   Eukaryota; Metazoa;
Mammalia; Eutheria;
SEQUENCE OF 1-22 |
MEDLINE=96032749;
                                                                                                  Submitted
                                                                                                                                                Yin L.Y., Wu Y., Patterson C.; "Full length human KDR/flk-1 s
"Submitted (DEC-1997) to the EM
                                                                                                                                                                                                                            KDR OR FLK1
                                  "Identification of a new endothelial cell growth factor receptor tyrosine kinase.";
                                                                                SEQUENCE OF 3-1356 FROM N.A.
                                                                                                                                 SEQUENCE FROM N.A.
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uman VEGF receptor KDR (VI
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Terman B.I., Dougher-Vermazen M., Carrion M.E., Dimitrov
Armellino D.C., Gospodarowicz D., Boehlen P.;
"Identification of the KDR tyrosine kinase as a receptor
                                                                                                 the European Bioinformatics Institute.
                                                                                                                 This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EN
                                                                                                                                                                                                                                                                                                                                                                            FUNCTION
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                                             an email to license@isb-sib.ch).
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CSF-1/PDGF
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DR PROSITE; PS00107; PROTEIN_KINASE_ATP; FALSE_NEG.
DR PROSITE; PS00109; PROTEIN_KINASE_TYR; 1.
DR PROSITE; PS00240; RECEPTOR_TYR_KIN_II; 1.
DR PROSITE; PS00240; RECEPTOR_TYR_KIN_II; 1.
DR PROSITE; PS50011; PROTEIN_KINASE_DOM; 1.
W Signal; Transferase; Tyrosine-protein kinase: Page 1 of the protein; Phosphorylation: Americal Signal.
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Search completed: June Job time: 569 sec
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Q28106 bos taurus
Q9w675 brachydanio
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Q63155 rattus norv
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Q92495 homo sapien
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ALIGNMENTS

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Ouery Match 89.1%; Score 57; DB 4; Length 374; Best Local Similarity 91.7%; Pred. No. 0.0032; Matches 11; Conservative 0; Mismatches 1; Indels 0; Gaps 0; 1 EDSGLYOCEAAT 12 1	Invest. 90:2102-2109(1992). 3418; AAA36049.1; - 2319; IALT. ; IPR003006; - 00047; 19; 3. PD002534; -; 1. 974 AA; 42632 MW; D33D59398CEEA699	SEQUENCE FROM N.A. SEQUENCE FROM N.A. TISSUE-BLOOD; MEDLINE-93055454; PubMed-1430234; C Porges A.J., Redecha P.B., 'Doebele R., Pan L.C., Salmon J.E., Kimberly R.D.; "Novel Fc gamma receptor I family gene products in human mononuclear	CEPTOR I. s (Human). Metazoa; Chordata; Cra utheria; Primates; Cat 9606;	Q92663; PRELIMINARY; PRT; 374 AA. Q92663; 01-FEB-1997 (TrEMBLrel. 02, Created) 01-FEB-1997 (TrEMBLrel. 02, Last sequence update) 01-OCT-2000 (TrEMBLrel. 15, Last annotation update)

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AC Q90
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Matches 11
     Giordano S., Laessing U., Lott
Submitted (APR-1996) to the EI
EMBL; U55211; AAA99159.1; --
HSSP; P20241; 1CFB.
INTERPRO; IPR001777; --
INTERPRO; IPR003006; --
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Q1-FEB-1997 (TremBLrel.
Q1-FEB-1997 (TremBLrel.
Q1-QCT-2000 (TremBLrel.
FC GAMMA RECEPTOR TYPE
CD64 OR FC<GAMMA>RIB.
                                                                                                                                                                                                                         Carassius auratus (Goldfish).
Eukaryota; Metazoa; Chordata; Craniata;
Actinopterygii; Neopterygii; Teleostei;
Cypriniformes; Cyprinidae; Cyprininae; Cypriniae; Cyprinidae; Cypriniae; Cypr
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Q90284;
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                                                                                                                                                                       SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                      L1-LIKE CELL ADHESION MOLECULE ANTIGEN
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Submitted (DEC-1992)
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Mammalia; Eutheria;
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S45704;
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1PR001777; -.
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(TrEMBLrel. 02, Last sequence update)
(TrEMBLrel. 15, Last annotation update)
CEPTOR TYPE I (FC GAMMA RIB-FC GAMMA RECEPTOR)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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PubMed=1402657;
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ne EMBL/GenBank/DDBJ
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Euteleostei;
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                                                                                                                                       Stuermer C.A.O.;
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Best Local
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Q90874;
01-NOV-1996
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PFAM; PF00047; ig
PRINTS; PR00014;
SEQUENCE 1232 A
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01-MAY-1999
01-MAY-1999
01-MAY-2000
SEQUENCE FROM N.A.

MEDLINE=92175529; PubMed=1531799;
Guida S., Heguy A., Melli M.;
"The chicken IL-1 receptor: differential evolution of the cytoplasmic
                                                                                                                                                                                                                                                                      IL-1 RECEPTOR I PRECURSO Gallus gallus (Chicken).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NON_TER
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Diaz M., Greenberg A.S., Flajnik M.F.;
"Somatic hypermutation of the new antigen receptor gene (NAR) nurse shark does not generate the repertoire: possible role in antigen-driven reactions in the absence of germinal centers.";
Proc. Natl. Acad. Sci. U.S.A. 95:14343-14348(1998).
                                                                                                                                                       NCBI_TaxID=9031;
                                                                                                                                                                                                              Archosauria;
                                                                                                                                                                                                                                          Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ginglymostoma cirratum (Nurse shark).
Eukaryota; Metazoa; Chordata; Craniata; V
Elasmobranchii; Galeomorphii; Galeoidea;
Ginglymostomatidae; Ginglymostoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PFAM; PF00047; ig;
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8; Conserv
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8; Conserv
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105 AA;
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(TrEMBLrel. 10,
(TrEMBLrel. 13,
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                                                                                                                                                                                                              Neognathae;
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11437
                                                                                                                                                                                                        Chordata; Craniata; Vertebrata; Eute
Neognathae; Galliformes; Phasianidae;
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66.7%;
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No.
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0.049;
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; Orectolobiformes;
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and extracellular domains."; Gene 111:239-243(1992). EMBL; M81846; AAA48924.1; -. HSSP; P14778; 1ITB.

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IPR003006;

IPR000157; -.

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01-NOV-1996
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01-AUG-1998 (TrEMBLrel. 07, Last seg
01-MAY-2000 (TrEMBLrel. 13, Last ann
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EMBL; D38492; BAA07504.1;'-.
HSSP; P40189; 1BQU.
                                                                                                                                                                                                                                                                                                             Takenawa T.,
                                                                                                                                                                                                                                                                                                                     Hosoya H
                                                                                                                                                                                                                                                                                                                                         STRAIN=WISTAR;
                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                  Rattus norvegicus (Rat).
                                                                                                                                                                                                                                                                                                                                                                                                          01-MAY-2000 (TrEMBLrel. 13, Last NEURAL ADHESION MOLECULE F3,
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                                                                                                                                                                                                                                                                                         rat brain.
                                                                                                                                                                                                                                                                                                                             STRAIN=WISTAR; TISSUE=BRAIN;
MEDLINE=95295987; PubMed=7777204;
                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=10116;
                                                                                                         303
                                                                                                                                                                                                                                                                                                  Developmental expression
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00047; 19; 6.

PS00092; N6_MTASE; UNKNOWN_1.

1021 AA; 113494 MW; FC8DC13055EE5C68
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                                                                                                                                                                                                                                                                                                                     Shimazaki K., Kobayashi
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                    (TrEMBLrel. 05, (TrEMBLrel. 08, (TrEMBLrel. 13,
                                                                                                                                               Conservative
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                                                           PRELIMINARY;
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IL-1 RECER
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Pred. No. 0.53;
1; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                             Craniata; Vertebrata; Sciurognathi; Muridae;
                                                          PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           L-1 RECEPTOR I.
4DB56DD7B1BB17AA CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                           COMPLETE
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No. 0.28;
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; Murinae; Rat
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1 EDSGLYQCEAAT 12

Query Match Best Local S Matches 7

Similarity 7; Conser

Conservative

73.4%;

Score 47; I Pred. No. 0.

DB 11; 0.32;

Length 412;

Indels

0,

Gaps

0,

Mismatches

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RESULT
Q63611
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Best Local S
Matches
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INTERPRO; IPR001777; -
INTERPRO; IPR003006; -
IPRAM; PF00041; fn3; 3;
PFAM; PF00047; ig; 5;
SEQUENCE 1256 AA; 1
                             Signal.
SIGNAL
                                                                                                                                STRAIN-BDIX; TISSUE-COLON TUMOR; MEDLINE-94253144; PubMed-8195207; Chadeneau C., LeMoullac B., Denis M.G.
                                                                                                                                                                                                                                                             01-NOV-1996 (TrEMBLrel. 01, Created 01-NOV-1996 (TrEMBLrel. 01, Last second-way-2000 (TrEMBLrel. 13, Last an TUMOR-ASSOCIATED ANTIGEN PRECURSOR.
                                                                                                                                                                                                                                                                                                      Q63611;
01-NOV-1996
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; l
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;
NCBI_TaxID-10116;
     SEQUENCE
                  CHAIN
                                                                                                                                                                                                             Rattus norvegicus (Rat).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                 Q63611
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Krauss R.S., Kang J.S., Submitted (SEP-1998) to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Krauss R.S., Kang J.S., Submitted (MAY-1997) to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Krauss R.S., Kang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A. MEDLINE=97362072; | Kang J.S., Gao M.,
                                                       PFAM; PF00047; ig;
                                                                                           J. Biol. Chem.
                                                                                                                                                                         SEQUENCE FROM N.A.
                                                                                                                                                                                                NCBI_TaxID=10116;
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                                                                    INTERPRO;
                                                                                                                     A novel member of the immunoglobulin
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                                                                                                       carcinoma cell lines.
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                                                                               L12025; AAB80767.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
9; Conserv
1
34
412
                                                                    IPR003006;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         gene-, serum-, and anchorage-regulated member of the n type III repeat family."; 138:203-213(1997).
     ΑĄ,
                                                                                             269:15601-15605(1994)
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, Feinleib J.L.;
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     MW;
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EMBL/GenBank/DDBJ
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EMBL/GenBank/DDBJ
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0; Mismatches
    POTENTIAL.
TUMOR-ASSOCIATED ANTIGEN.
98025D4C2161859B CRC64;
                                                                                                                                                                                                              Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Rat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 775805754F0C2EA4 CRC64;
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                                                                                                                                                                                                                                                                                                                                 412 AA
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                                                                                                                     superfamily expressed
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; Murinae; Rattus
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Best Local
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Q60977;
Q1-NOV-1996
                                                      immunoglobulin superfamily.";
Mamm. Genome 7:636-637(1996).
EMBL; U35836; AAB17502.1; -.
HSSP: Q13740; 1KJC.
                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                             Chadeneau C., LeMoullac B., LeCabellec
                                                                                                                                                                                                                                                                                                                                                                                   TUMOR-ASSOCIATED GLYCOPROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                          01-OCT-2000
                                                                                                                                                                                                                             STRAIN=BALB/C
                                                                                                                                                                                                                                                                                                                                                                  TAA1 OR TAGE4.
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                                                                                                                                                                  Denis M.G.
                                                                                                                                                                                                    TEDLINE-96327659; PubMed=8678999;
                                                                                                                                                                                                                                            SEQUENCE FROM: N.A.
                                                                                                                                                                                                                                                                                    NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                               Mus musculus (Mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Submitted
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"Structure of the rat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-MAY-2000 (TrEMBLrel. 01-OCT-2000 (TrEMBLrel.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRAIN=BN/SSNHSD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TUMOR-ASSOCIATED GLYCOPROTEIN
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                                                                                                                                       Isolation and chromosomal location
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                MGI:107741; Taal.
RPRO; IPR001064; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EDEGIYECQIAT 134
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EDEGIYECQIAT 134
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IPR003006; -
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Rodentia;
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Rodentia;
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01,
15,
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EMBL/GenBank/DDBJ databases
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Pred. No.
                                                                                                                                                                                                                                                                                             Craniata; Vertebrata; Euteleostomi;
Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                   E4.
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RESULT
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AC Q6
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Best Local
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INTERPRO; IPR003006; -.
PFAM; PF00041; fn3; 6.
PFAM; PF00047; -14; 6.
SEQUENCE 1287 AA; 144
Q63155 PRELIMINARY;
Q63155;
01-NOV-1996 (TremBirel. 01,
                                                                                                                                                                                                                                                                                                                                              STRAIN-BRISTOL N2;
Du Z., Gattung S.;
"The sequence of C. e
Submitted (MAR-1996)
                                                                                                                                                                                                               EMBL; U50067; AAA33439.2; -. HSSP; P20241; 1CFB.
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01-NOV-1996
01-MAY-2000
01-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
STRAIN=BRISTOL N2;
MEDLINE=99069613; PubMed=9851916;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Caenorhabditis elegans.
Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida; Rhabditoidea; Rhabditidae; Peloderinae; Caenorhabditis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PROSITE;
SEQUENCE
                                                                           168
                                                                                                                                                                                                                                                            STRAIN-BRISTOL N2;
                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A. STRAIN-BRISTOL N2;
                                                                                                                                                                                                                                                                                                                                                                                                        investigating biology. The C
Science 282:2012-2018(1998).
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                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                       Vaterston
                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=6239;
                                                                                                                                                                                                                                                                                              Submitted
                                                                                                                                                                                                                                                                                                                                                                                                                             Genome sequence of the nematode C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 123 EDEGIYECQIAT 134
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                                                                         DAGLYOCEA 176
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                                                                                                                   Similarity
8; Conserv
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7; Conserv
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415 AA; 45280 MW; 159AF8D60EF90D0F CRC64;
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Matches 8
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01-OCT-2000
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novel receptor expressed on neutrophils and monocytes.";
J. Immunol. 164:4991-4995(2000).
EMBL; AF7287008; AAF90197.1; -.
Receptor.
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Culotti J.G., Tessier-Lavigne
"Deleted in Colorectal Cancer
Cell 87:175-185(1996).
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
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EMBL; M32291; AAA41086.1;
HSSP; P56276; ITLK
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                                                                                                                                               SEQUENCE FROM N.A. MEDLINE=20261626; PubMed=10799849;
                                                                                                                                                                                                                                                          Gingras M.C., Lapillonne H. Monocytic differentiation
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Mammalia; Eutheria;
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                                                                                                      Bouchon A., Dietrich J., Colonna M.; "Cutting edge: inflammatory responses
                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=9606;
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Metazoa; Chordata; C
Metazoa; Primates; (
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S.R., Preisinger
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                                                                                                                                                                                                                                                                                                                                                                     Craniata; Vertebrata; Catarrhini; Hominidae;
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; Murinae; Rattus
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G., Kinzler
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Q28106;
                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Actinopterygii; Neopterygii; Teleostei; Euteleostei;
Cypriniformes; Cyprinidae; Rasborinae; Danio.
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                       Warren J.T.
Kuwada J.Y.;
                                                                                                                                                                                      Brachydanio rerio (Zebrafish) (Zebra danio)
                                                                                                                                                                                                                              TRANSIENTLY
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"Cloning of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mammalia; Eutheric
Bovidae; Bovinae;
                                                                SEQUENCE FROM N.A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TISSUE-BRAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PFAM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=9913;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INTERPRO;
'Molecular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  106 EDSGLYQC 113
                                                                                                                                                                                                                                                                                                                                                                                                                                        303
                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 EDSGLYQCEA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EDSGLYQC 8
                                                                                                                                                                                                                                                                                                                                                                                                                                        EDEGIYECEA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PF00047; ig; 6.
TE; PS00092; N6_MTASE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    IPR003006;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       IPR002052;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             IPR001777;
  cloning
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (Bovine)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1018 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      234 AA;
                                        Jr.,
                                                                                                                                                                                                                                             (TrEMBLrel. 12,
(TrEMBLrel. 12,
(TrEMBLrel. 13,
                                                                                                                                                                                                                              EXPRESSED AXONAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            the cDNA
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                                                                                                                                                                                                                                                                                                                                   PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Shimazaki K.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BAA06861.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                        312
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10
                                        Chandrasekhar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bos
  and developmental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chordata; Craniata; Vertebrata; Euteleostomi; Cetartiodactyla; Ruminantia; Pecora; Bovoidea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26386 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20 POTENTIAL.
113384 MW; 551FC10ED7920341 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        encoding neural
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       71.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          . 9%;
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                                                                                                                                                                                                                                                  Created)
Last sequence update)
Last annotation updat
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Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 46;
Pred. No.
                                                                                                                                                                                                                              GLYCOPROTEIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 46; DB
Pred. No. 1.3;
                                        Α.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AA114696E35D4D45 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                        Kanki J.P.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1018
                                                                                                                                                                                                                                                                                                                                   1040
  expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Fukamauchi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0 B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       в 4;
. 28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              6;
                                                                                                                                                                                                                                             update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                        Rangarajan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 1018,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length
  analysis of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                  Euteleostomi
                                                                                                                                             Ostariophysi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Takenawa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bovoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
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RT zebrafish axonal glycoprotein tag1.";

RL Submitted (MAY-1998) to the EMBL/GenBank/DDBJ databases.

DR EMBL; AF064799; AAD31083:1; -.

DR INTERPRO; IPR001777; -.

DR PFAM; PF000041; fin3; 4.

DR PFAM; PF000047; ig; 6.

SQ SEQUENCE 1040 AA; 114626 MW; E6C2DB8B177D1B3C CRC64;

``

GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.

```
Title:
Perfect score:
Sequence:
 Result
No.
 Database
 Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
 Minimum DB seq length: 0
Maximum DB seq length: 200000000
 Run on:
 OM protein -
 Total number of hits satisfying chosen parameters:
 Searched:
 Scoring table:
 Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
 Score
 protein search, using sw model
 Query
 100.0
 BLOSUM62
Gapop 10.0 , Gapext 0.5
 June 4, 2001, 12:13:31; Search time 106.14 Seconds (without alignments)
 390729 seqs, 57163235 residues
 us-09-284-107-27
61
 A_Geneseq_0401:*
 1 EDGNVLKRSPEL 12
 Length DB
 12
261
344
344
 IJ
 W60559
 R22549
 /geneseq/geneseqp/AA1
 SUMMARIES
 'geneseqp/AA'
 'qeneseqp,
 (without alignments)
6.463 Million cell updates/sec
 390729
 995.DAT:
996.DAT:
997.DAT:
 Human macrophage-s
Human FCRI (CDNA c
Human macrophage-s
Human macrophage-s
Human FCRI (CDNA C
 Description
 Oligopeptide from
 sfc-gammaRI
Fc receptor
 Fc receptor
 receptor
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|                                        |        |     | -,    |       | Ý. |  |
|----------------------------------------|--------|-----|-------|-------|----|--|
| Arabidopsis thalia                     | 316    | 21  | 156   | 57.4  |    |  |
| reted pr                               | G02088 | 21  | 89    | 57.4  | 35 |  |
| dopsis                                 | 315    | 21  | 586   | 59.0  |    |  |
| a func                                 | 977    | 18  | . 507 |       |    |  |
| etec                                   | 74     | 21  | 359   | •     |    |  |
| 9                                      | B44246 | 21  | 359   | •     |    |  |
| _                                      | Y41690 | 20  | 359   |       |    |  |
| Human cell surface                     | Y94336 | 21  | 358   | •     |    |  |
| bidopsi                                | G42149 | 21  | 321   | •     |    |  |
| rF gene produ                          | u      | 17  | 288   |       |    |  |
| S. pneumoniae deri                     | Y86103 | 19  | 132   | 59.0  |    |  |
| is thal                                | 5      | 21  | 753   | 60.7  |    |  |
| Arabidopsis thalia                     | 5      | 21  | 715   | 60.7  |    |  |
| Arabidopsis thalia                     | в      | 21  | 673   | 60.7  |    |  |
| Arabidopsis thalia                     | œ      | 21  | 637   | 60.7  |    |  |
| <ul> <li>Arabidopsis thalia</li> </ul> | G60546 | 21  | 268   | 60.7  |    |  |
| Arabidopsis thalia                     | G58313 | ·21 | 268   | 60.7  |    |  |
| Arabidopsis thalia                     | G60547 | 21  | 261   | 60.7  |    |  |
| Arabidopsis thalia                     | G58314 | 21  | 261   | 60.7  |    |  |
| idopsis thal                           | G60548 | 21  | 140   | •     |    |  |
| Arabidopsis thalia                     | G58315 | 21  | 140   | ٠     |    |  |
| is thal                                | G13324 | 21  | 523   | •     |    |  |
| Arabidopsis thalia                     | G13325 | 21  | 483   | •     |    |  |
| is                                     | G13326 | 21  | 431   | 62.3  |    |  |
| įst                                    | G28824 | 21  | 421   | •     |    |  |
| is thal                                | G28825 | 21  | 407   | •     |    |  |
| is thali                               | G12849 | 21  | 163   | •     |    |  |
| idopsi                                 | G12850 | 21  | 149   | •     |    |  |
| id Fc                                  | R12428 | 12  | 410   | •     |    |  |
| Human Fc receptor                      | 619    | 20  | 377   | •     |    |  |
| n macrophage-                          | 9613   | 21  | 374   | 86.9  |    |  |
| macr                                   | 9618   | 21  | 344   | 86.   |    |  |
| er assc                                | 368    | 21  | 399   |       |    |  |
| Human high affinit                     | Y96226 | 21  | 374   | 100.0 |    |  |

## ALIGNMENTS

RESULT W60559

W60559 standard; peptide; 12 AA

SAXAXE

W60559;

18-AUG-1998 (first entry)

Oligopeptide from extracellular domain of CD64.

| _                     |                                                                     |                                                                    |                                                                    | _  |                      |    |                            | _  | _                          |   |                            |    | _                          |    |              | _    |               |    |           |    | _                                              | _                                                           |                                                       |  |
|-----------------------|---------------------------------------------------------------------|--------------------------------------------------------------------|--------------------------------------------------------------------|----|----------------------|----|----------------------------|----|----------------------------|---|----------------------------|----|----------------------------|----|--------------|------|---------------|----|-----------|----|------------------------------------------------|-------------------------------------------------------------|-------------------------------------------------------|--|
| PT.                   | PΤ                                                                  | ΡŢ                                                                 | PΤ                                                                 | XX | DR                   | XX | Ιď                         | XX | PA                         | · | PR                         | XX | Э₽                         | XX | PD           | · XX | ,<br>PN       | XX | SO        | XX | KW                                             | KW                                                          | KW                                                    |  |
| specific antibodies / | oligo:peptide derived from the target, useful for, e.g. identifying | expressing on phage surface and testing for binding to immobilised | Identifying peptide(s) binding specifically to protein target - by |    | WPI; 1998-240964/21. |    | De Kruif CA, Logtenberg T; |    | (UYUT-) RIJKSUNIV UTRECHT. |   | 08-OCT-1996; 96EP-0202791. |    | 07-OCT-1997; 97WO-NL00557. |    | 16-APR-1998. |      | WO9815833-A1. |    | Synthetic |    | cell transformation; mutation; anti; oncogene. | immunohistochemical; immunofluorescent analysis; detection; | Extracellular domain; CD64; identification; antibody; |  |

Example 1; Page 29;

40pp; English

```
DI DR RESERVE PO DE PRESERVE PO DE P
 QΥ
 밁
 Query Match
Best Local Similarity
 Baell JB,
Maxwell K
 11-SEP-1998;
06-FEB-1998;
 For receptor; Fc-gammaRIIa; human; FCR; model; three-dimension; 3-D; atomic coordinate; bioactive compound design; computer-assisted; drug design; therapy; inhibitor; Fc-gammaR; Fc-epsilonR; IgG; IgE; tissue damage; hypersensitivity; inflammatory cell recruitment; inflammatory modulator; Fc-gammaRIa; immune function regulation;
 (ILEX-) ILEXUS PTY LTD.
 04-FEB-1999;
 12-AUG-1999.
 WO9940117-A1
 Homo sapiens
 inflammatory modula
anti-inflammatory;
 15-NOV-1999
 Y33183
 surface of a replicable display package, synthesising oligopeptides derived from the target protein on a solid phase, and testing for binding between the peptide and oligopeptides. The method is used to screen large peptide libraries, especially to detect antibodies, or their fragments, that bind to cell markers or that can differentiate between different forms of the same protein, including bispecific antibodies that bind to epitopes on the same monometric antigen or two epitopes on different molecules. The genes/oligonucleotides that encode selected peptides can be isolated and used for recombinant production of antibodies. Selected antibodies are used, e.g. for immunohistochemical or
 Y33183 standard;
 Sequence
 Synthetic oligopeptides W60537-61 are derived from the extracellular domain of CD64. They were synthesised on 25 polyethylene rods as adjacent, non-overlapping sets. Peptide specific phages were isolated by affinity binding to the CD64-covered rods. The specification describes a method for the identification of a peptide able to bind specifically to a target protein. The method comprises displaying the peptide on the surface of a renlicable displaying the peptide on the
 by mutation in (anti)oncogenes.
 Synthetic oligor domain of CD64.
 antibodies. Selected antibodies are used, e.g. for immunohistochemical or immunofluorescent analysis, and also to detect cell transformation caused
 1 edgnvlkrspel 12
 1 EDGNVLKRSPEL 12
 KF,
 Epa V, Garrett TPJ, McKe, McCarthy TD; McKe
 12
 100.0%;
ilarity 100.0%;
Conservative (
 (first entry)
 ΑĄ;
 98US-0099994.
98US-0073972.
 99WO-IB00367.
 Protein;
 protein fragment.
 immunoprotective; sfc-gammaRI.
 261
 McKenzie IFC,
 0;
 A
 Score 61; DB 19;
Pred. No. 3.4e-05;
 Mismatches
 PM, Matthews Pietersz GA;
 Length 12;
 Indels
 BR;
 0;
```

Three-dimensional structures and models of Fc receptors, useful in computer-assisted drug design

This amino acid sequence was predicted from the cDNA sequence of cDNA clone p98/X2. It differs from the sequence predicted from clone p135 (see Q21178) at position 25; p135 encodes a Ser residu while clone p98/X2 predicts a Thr residue. At position 58, p90 '

residue p90

Example 10;

Page 94a;

160pp; English.

neoplasms,

etc.

WPI; 1999-539978/45

```
CCC CCC XXX
 PD
 PN XX
 Query Match
Best Local Similarity
---hes 12; Conserv
 В
 Qγ
 RESULT
 Seed
 New CD53 cell surface antigen and DNA encoding immuno-therapy and diagnosis of haematopoietic
 N-PSDB; Q21179.
 reduce IgG- or IgE-mediated tissue damage, e.g. IgG- or IgE-mediated hypersensitivity, recruitment of inflammatory cells or release of inflammatory modulators. The therapeutic compositions can also be used to treat disease in animals. The knowledge of the 3-D structure of Fc-gamma compounds that regulate immune function and inflammation in an animal, including humans (i.e. structure based drug design). For example, chemical compounds can be designed to block binding of immunoglobulin to an Fc receptor protein using various computer programs and models. The products of the invention have anti-inflammatory and immunoprotective activity. This sequence represents the human sFC-gammaRI protein fragment described in the method of the invention.
 WPI; 1992-056864/07
 (GEHO-) GEN HOSPITAL CORP.
 13-JUL-1990;
 15-JUL-1990; '
 23-JAN-1992
 WO9201049-A
 Homo sapiens
 Rapid immunoselection cloning technique; cell surface antigen, immunodiagnosis; high affinity receptor.
 Human macrophage-specific FcRI receptor encoded by clone p98.
 21-MAY-1992
 R22549 standard; Protein;
 oligonucleotides, peptides, peptidomimetics and small organic molecules by computer-assisted methods of drug design. Therapeutic compositions that inhibit the activity of Fc-gammaR or Fc-epsilonR can be used to
 Sequence 261 AA;
 the FCR can be used to design bloactive chemical compounds,
 representing a three-dimensional (3-D) structure that substantially conforms to the specified atomic coordinates. Computer model images of
 This invention describes a novel model of an Fc receptor (FCR) protein representing a three-dimensional (3-D) structure that substantially
 Claim 11; Page 316-317; 326pp; English
 246 edgnvlkrspel 257
 1 EDGNVLKRSPEL 12
 ω
 В,
 Aruffo A,
 100.0%; ilarity 100.0%; Conservative 0
 90US-0553759
 90WO-US04986
 Amiot M;
 0;
 Score 61; DB Pred. No. 0.0; Mismatches
 0.001;
 DB 20;
 ŗ.
 0;
 Length 261;
 for
 0,
 Gaps
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0;

888888888<del>x</del>&

å 밁

0;

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RESULT
R91439
 Query Match
Best Local
 Matches
 The amino acid sequence (R91439) of human FCRI was detd. from a CDNA clone, p98 (T14718), obtd. from a CDNA library using an immunoselection cloning method. FCRI is a high affinity receptor for the Fc portion of IgG, normally located on cell surfaces of macrophages. 2 Other cDNA clones, p135 (T14717) and p90 (T14719), macrophages.
 01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
 WPI; 1996-200279/20
N-PSDB; T14718.
 and
 (see Q21180) predicts Val and p98/X2 predicts Leu. Sequences predicted from all 3 clones show the typical features of a type integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane-spanning doma
 Aruffo
 13-JUL-1990;
18-OCT-1993;
 Misc-difference
 Homo sapiens
 therapy; diagnosis;
 Cell surface antigen; cloning; immunoselection; immunotherapy;
 Human FcRI (cDNA clone p98 product).
 R91439;
 R91439 standard; Protein;
 Sequence
 cys
 Example 10; Column 55-56; 79pp; English.
 25-FEB-1988;
 09-APR-1996
 US5506126-A
 30-OCT-1996
 portion
 (GEHO) GEN HOSPITAL CORP
 ortion contains six potential N-linked glycosylation sites and six ys residues distributed among three C2 set 1g-related domains. fusion protein of FCRI and a receptor ligand will be helpful to ncrease the potency of antibodies in therapy.
 1 EDGNVLKRSPEL 12
the p98 product
 a short, highly charged cytoplasmic domain. The extracellular tion contains six potential N-linked glycosylation sites and si
 12;
 Similarity
 344 AA;
 Conservative
 (first entry)
 92US-0983647.
88US-0160416.
89US-0379076.
 900S-0553759
930S-0139273
 88US-0160416
 В;
 Location/Qualifiers 58
 /note=
 vector;
 100.0%; Score 61;
100.0%; Pred. No.
tive 0; Mismatc
 truncated compared
 "amino acid 58 is Leu in clone translated product"
 344
 FCRI; Fc receptor
 A
A
 Mismatches
 0.0014;
with those of the p135
 DВ
 13;
 useful for isolation
 Length 344;
 Indels
 p135
 domain,
 0;
 0
```

Query Match
Best Local Similarity
Matches 12; Conser

Conservative

0;

100.0%;

Score 61; Pred. No.

Length 374;

Indels

0,:

Gaps

0

DB 13; Lu. No. 0.0016; Mismatches

Sequence

374

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ភភពនុង
ភភពនុង
В
 QY
 RESULT
 IJ
 Best Loc
Matches
 Query Match
 This amino acid sequence was predicted from the cDNA sequence of cDNA clone pl35. It differs from the sequence predicted from clones p98/X2 and p90 (see Q21179 and Q21180, respectively) at position 25; pl35 encodes a Ser residue and the other two clones predict a Thr residue. At position 58, pl35 predicts Leu and p90 predicts Val. Sequences predicted from all 3 clones show the
 . Sequence
 and p90 products (see also R91438 and of FCRI and a receptor ligand will be potency of antibodies in therapy.
 distributed among three C2 set Ig-related domains of FCRI and a receptor ligand will be helpful to i potency of antibodies in therapy.
 hydrophobic membrane-spanning domain, and a short, highly cytoplasmic domain. The extracellular portion contains six
 typical features of a type I integral membrane protein and a short hydrophobic signal sequence, a single 21-residue
 R20811;
 R20811 standard;
 WPI; 1992-056864/07.
 Seed B,
 15-JUL-1990;
 Rapid immunoselection
 Human macrophage-specific FcRI receptor encoded by clone p135
 21-MAY-1992
 potential N-linked glycosylation sites and six Cys residues
 Example
 New CD53 cell surface antigen and DNA encoding immuno-therapy and diagnosis of haematopoietic
 N-PSDB; Q21178
 (GEHO-) GEN
 13-JUL-1990;
 23-JAN-1992
 WO9201049-A
 Homo sapiens
 265 edgnvlkrspel 276
 Local
 1 EDGNVLKRSPEL 12
 σ
 ch 100.0%; l Similarity 100.0%; 12; Conservative (
 10;
 344 AA;
 HOSPITAL CORP
 Page
 (first entry)
 90US-0553759
 90WO-US04986
 94a;
 Protein;
 cloning technique;
affinity receptor.
 160pp;
 374 AA
 0;
 English.
 Score 61; DB
Pred. No. 0.0
); Mismatches
 W00859).
 helpful
 0.0014;
 cell surface antigen;
 DB 17;
 . A fusion protein in increasing the
 0
 it -
 neoplasms,
 increase the
 Length 344;
 Indels
 A fusion
 highly charged
 for
 etc.
 include
 protein
 0
 Gaps
```

B Š

265.

1 EDGNVLKRSPEL 12 |||||||||||| 65 edgnvlkrspel 276

```
RESULT
R91438
ID RS
43 X X X
 밁
 Ş
 Query Match
Best Local Similarity
Matches 12; Conser
 cDNA clone p90. It differs from the sequence predicted from clone p135 (see Q21178) at position 25, p135 encodes a Ser residue and clone p90 predicts after residue. At position 58, p90 predicts val while clones p135 and p99/X2 (see Q21179) both predict Leu. Sequences predicted from all 3 clones show the typical features of a type I integral membrane protein and include a short hydrophobic domain, and a short, highly charged cytoplasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and six Cys residues distributed among three C2 set Ig-related domains. A fusion protein of FCRT and a receptor ligand will be helpful to increase the potency of antibodies in therapy.
 30-OCT-1996
 R91438;
 R91438 standard;
 Sequence
 This amino acid sequence was predicted from the cDNA sequence cDNA clone p90. It differs from the sequence predicted from clone p135 (see Q21178) at position 25; p135 encodes a Ser res
 265
 Example 10; Page 94a; 160pp; English.
 New CD53 cell surface antigen and DNA encoding it for immuno-therapy and diagnosis of haematopoletic neoplasms, etc.
 WPI; 1992-056864/07
N-PSDB; Q21180.
 Seed B,
 13-JUL-1990;
 WO9201049-A
 (GEHO-)
 15-JUL-1990;
 Homo sapiens
 Human macrophage-specific FcRI receptor encoded by clone p90.
 R22550
 Rapid immunoselection cloning technique; cell surface antigen; immunodiagnosis; high affinity receptor.
 21-MAY-1992 (first entry)
 R22550 standard; Protein; 374
 1 EDGNVLKRSPEL 12
 edgnvlkrspel 276
 6
 GEN HOSPITAL CORP
 Aruffo A,
 374 AA;
 100.0%;
ilarity 100.0%;
Conservative
(first entry)
 90US-0553759
 90WO-US04986.
 Protein;
 Amiot M;
 374 AA.
 0
 Score 61; DB 13;
Pred. No. 0.0016;
Mismatches 0;
 AA.
 0;
 Length 374;
 Indels
 0;
 Gaps
 0;
```

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RESULT
W00859
XX DE XXX
 В
 ρ
 Query Match
Best Local Similarity 100
Matches 12; Conservative
 W00859;
 W00859 standard; Protein; 374 AA
 13-JUL-1989;
13-JUL-1990;
18-OCT-1993;
 Sequence
 The amino acid sequence (R91438) of human FCRI was detd. from a CDNA clone, pl35 (T14717), obtd. from a cDNA library using an immunoselection cloning method. FCRI is a high affinity receptor for the FC portion of IgG, normally located on cell surfaces of macrophages. Another isolated cDNA clone, p90 (T14719), coded for a variant (W00859) of the FCRI sequence, and a third clone, p90 (T14718), coded for an FCRI (R91439) having a different C-terminal sequence. A fusion protein of FCRI and a receptor ligand will be helpful in increasing the potency of antibodies in therapy.
 The amino acid sequence (R91438) of human FCRI was CDNA clone, p135 (T14717), obtd. from a cDNA librar
 Cloning of cDNA encoding cell surface antigen - useful for isolation of diagnostic and therapeutic proteins {\bf r}
 Example 10; Column 55-56; 79pp; English.
 N-PSDB;
 WPI; 1996-200279/20
N-PSDB; T14717.
 Aruffo A,
 01-DEC-1992;
25-FEB-1988;
 Misc-difference
 25-FEB-1988
 09-APR-1996
 US5506126-A
 Cell surface antigen; cloning; immunoselection; immunotherapy; therapy; diagnosis; vector; FcRI; Fc receptor.
 (GEHO) GEN HOSPITAL CORP
 Misc-difference
 Homo sapiens.
 Human FcRI (cDNA clone p135 product)
 374 AA;
 Seed B;
 9305-0139273
 89US-0379076
90US-0553759
 88US-016041
 92US-0983647
 88US-0160416
 58
 Location/Qualifiers
 /note= "amino acid 58 is Val in p90 clone
 translated product"
 /note= "amino acid
translated
 100.0%;
 0
 Score 61;
Pred. No.
 Mismatches
 25 is Thr
 0.0016;
 DB 17;
 in clone p90 and
 Length 374;
 Indels
 .
0;
 p98
 Gaps
 P98
```

Cell surface antigen; cloning; immunoselection; immunotherapy;

Human FcRI (cDNA clone p90 product).

30-OCT-1996

(first entry)

0,

```
8 x 0 0 0 0 0 0 0 0 x 8
 Query Match
Best Local Similarity
Matches 12; Conser
 FCRI and a receptor li
potency of antibodies
 The amino acid sequence (W00859) of human FCRI was detd. from a CDNA clone, p90 (T14719), obtd. from a cDNA library using an
 Cloning of cDNA encoding cell surface antigen - useful for isolation of diagnostic and therapeutic proteins
 N-PSDB;
 Aruffo A.
 Misc-difference
 Misc-difference
 Misc-difference
 cloning.
 07-JUN-1999
 W80448;
 cDNA clone, p90 (T14719), obtd. from a cDNA library using an immunoselection cloning method. FCRI is a high affinity recept for the Fc portion of IgG, normally located on cell surfaces of macrophages. Other cDNA clones (see also T14717-18) coded for variants (R91438-39) of the FCRI sequence. A fusion protein of FCRI and a receptor ligand will be helpful in increasing the
 WPI; 1996-200279/20
 09-APR-1996.
 US5506126-A.
 Misc-difference
 Homo sapiens
 Fc receptor I;
 Human Fc receptor I.
 W80448 standard; Protein; 374 AA
 Example 10; Column 55-56; 79pp; English.
 (GEHO) GEN HOSPITAL CORP
 18-OCT-1993;
 25-FEB-1988;
 01-DEC-1992;
 25-FEB-1988;
 Homo sapiens
 therapy; diagnosis; vector; FcRI; Fc receptor.
 13-JUL-1989;
13-JUL-1990;
 265 edgnvlkrspel 276
 1 EDGNVLKRSPEL 12
 T14719.
 374
 Conservative
 (first entry)
 92US-0983647.
88US-0160416.
89US-0379076.
90US-0553759.
93US-0139273.
 A,
 FCRI;
 88US-0160416
 23
/note= "encoded by CTG
 Location/Qualifiers
 'note=
 'note=
 note=
 100.0%;
 cell surface antigen; lymphocyte;
 in therapy
 "encoded by GAC"
 "encoded by TGG"
 "encoded by CTC"
 0;
 Score 61;
Pred. No.
 Mismatches
 FCRI is a high affinity receptor llv located on cell surfaces of
 0.0016;
 0,
 Length 374;
 Indels
 0
 Gaps
```

```
1.4
1.4
1.4
 antigens from mammalian lymphocytes (see W80440-55). The isolated genes can be expressed in a prokaryotic or eukaryotic host cells to produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V63444) which allow the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and
 New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially human lymphocyte antigenic sequences
 Misc-difference
 transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V63442–63) encoding cell surface antigens from mammalian lymphocytes (see W80440–55). The isolated
 adhesion to an antibody-coated substrate. The method is useful for the isolation and cloning of any protein which can be expressed and
 expression libraries is based on transient expression of an antigen by in eukaryotic cells and selection of cells expressing the antigen by
 were also obtained.
 This is the amino acid sequence of human Fc receptor I (FCRI), as deduced from cDNA clone p135 (see v63456) isolated using a rapid immunoselection cloning method from a cDNA library expressed in cells. The cDNA library was constructed from polyA RNA of cells
 WPI; 1998-609251/51
N-PSDB; V63456.
 Aruffo A,
 01-DEC-1992;
25-FEB-1988;
 21-MAY-1997;
 03-NOV-1998
 US5830731-A
 Misc-difference
 Misc;difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 from a single patient undergoing extracorporeal
 Example 10; Column 53.54; 75pp; English
 (GEHO) GEN HOSPITAL CORP.
 21-MAY-1997
 13-JUL-1989
13-JUL-1990
 mmunotherapeutic applications,
 including
 of
 92US-0983647.
88US-0160416.
89US-0379076.
90US-0553759.
 /note=
77
 /note=
256
 97US-0861205
 97US-0861205
 /note=
171
 /note=
99
 /note=
85
 176
 141
 103
 60
 /note=
 /note=
 'note-
 note-
 note=
 humans
 Clones p90
 A novel
 "encoded
 "encoded by
 *encoded by
 "encoded by
 "encoded by
 "encoded
 "encoded
 "encoded
 "encoded
 "encoded
 90 (see x07372) and p98 (see x07373) method for cloning cDNAs from mamma
 infections,
 useful for immunodiagnostic and including the diagnosis and fections, diseases, and disorder:
 ьy
 þу
 bу
 Ьy
 bу
 þу
 GGG
 GTC"
 ATG'
 AAC"
 GGC"
 CCC
 CAA"
 TCC"
 AAT"
 interleukin-
 from mammalian
 disorders
 FCRI), as
| a rapid
|ssed in CC
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Sequence

of

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100.0%;

В ΩV

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RESULT 10
W97833
 Best Local Similarity 100 Matches 12; Conservative
New cloning vector and polylinker - based on existing sequences
 WPI; 1998-609251/51
N-PSDB; X07372.
 Aruffo A,
 (GEHO) GEN HOSPITAL CORP
 13-JUL-1990;
21-MAY-1997;
 01-DEC-1992;
 21-MAY-1997;
 03-NOV-1998
 Misc-difference
 US5830731-A
 Misc-difference
 Misc-difference
 Misc-difference
 3-JUL-1989
 25-FEB-1988;
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Homo sapiens
 Fc receptor I; FcRI; cell surface antigen; lymphocyte;
 07-JUN-1999
 W97833;
 W97833 standard; Protein; 374 AA
 Human Fc receptor
 1 EDGNVLKRSPEL 12
||||||||||||
| 265 edgnvlkrspel 276
 Seed B;
 (first entry)
 92US-0983647.
88US-0160416.
89US-0379076.
99US-0553759.
97US-0861205.
 97us-0861205
 /note=
.171
 159
 /note=
256
 /note=
176
 /note= "encoded by CAA"
 /note=
99
 /note=
85
 /note=.
77
 60
 /note=
 /note= "encoded by
 Location/Qualifiers
 /note=
 note=
 'note=
 note- "encoded by
 "encoded
 "encoded by
 "encoded by
 "encoded by GTC"
 "encoded by GGC
 "encoded
 "encoded by
 "encoded by CTC"
 "encoded
 "encoded by ccc
 "encoded
 0
 Pred. No. 0.0016;
; Mismatches 0;
 bу
 by TCC
 γď
 by AAT"
 AAC"
 ATG"
 CCC.
 CTG"
 GAC"
 TGG"
 Indels
 human;
 0;
 for
 Gaps
 0;
RESULT
W97834
 Dp.
 Qy
 888
 Query Match
Best Local Similarity
Matches 12; Conser
 Misc-difference
 Misc-difference
 07-JUN-1999
 Homo sapiens
 W97834
 Sequence
 265
 11
```

0;

```
genes can be expressed in a prokaryotic or entaryotic host cells to produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V6344) which allow the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and immunotherapeutic applications, including the diagnosts and treatment of immune-mediated terminations.
 transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V63442-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated genes can be expressed in a prokarvotic or antigens.
 immunotherapeutic applications, includir treatment of immune-mediated infections,
 were also obtained. A novel method for cloning cDNAs from mammalian expression libraries is based on transient expression of an antigen by the antigen by
 in eukaryotic cells and selection of cells expressing adhesion to an antibody-coated substrate. The method the isolation and cloning of any protein which can be
 were also obtained.
 efficient clohing and expression of mammalian cDNA(s), especially human lymphocyte antigenic sequences
 from a single patient undergoing extracorporeal induction therapy. Clones plas (see V63456) an
 This is the amino acid sequence of human deduced from cDNA clone p90 (see X07372) immunoselection cloning method from a cDN
 Example 10; Column 53-54; 75pp; English
 The cDNA library was constructed from polyA RNA of cells
 including humans.
374 AA;
 lones p135 (see V63456) and p98 (see X07373) A novel method for cloning cDNAs from mammal
 cDNA library expressed
 Fc receptor I isolated using
 interleukin-2
 I (FCRI), .
ing a rapid
 is useful
 cos
```

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Misc-difference
 Misc-difference
 Misc-difference
 Human Fc receptor
 Misc-difference
 Misc-difference
 Fc receptor I;
 W97834 standard; Protein; 374
 1 EDGNVLKRSPEL
 edgnvlkrspel
 100.0%; ilarity 100.0%; Conservative
 (first entry)
/note=
. 77
 /note=
 FCRI;
 /note=
 45
 /note=
23
 Location/Qualifiers
 /note=
 276
 12
 cell surface antigen; lymphocyte; human;
 "encoded by
 "encoded by
 "encoded by TGG'
 "encoded by CTG"
 "encoded by
 0;
 Ą.
 Score 61; DB 19; Pred. No. 0.0016; Mismatches 0
 CCC.
 GAC"
 CTC"
 0;
 Length 374;
 Indels
 0;
 Gaps
```

tor To

Misc-difference

/note= /note= 103

"encoded by CAA" "encoded by TCC" /note= 85

"encoded

bу AAT

```
밁
 Matches
 Query Match
Best Local'
 expression libraries is based on transient expression of an antigen in eukaryotic cells and selection of cells expressing the antigen by adhesion to an antibody-coated substrate. The method is useful for the isolation and cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V63442-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated genes can be expressed in a prokaryotic or eukaryotic host cells to produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V6344) which allow the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and
 25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
21-MAY-1997;
 This is the amino acid sequence of human Fc, receptor I (FcRI), as deduced from cDNA clone p98/X2 (see X07373) isolated using a rapid immunoselection cloning method from a cDNA library expressed in COS cells. The cDNA library was constructed from polyA RNA of cells
 New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially human lymphocyte antigenic sequences
 WPI; 1998-609251/51.
N-PSDB; X07373.
 21-MAY-1997; \
 03-NOV-1998
 Misc-difference
 Misc-difference
 Misc-difference
 Sequence
 induction therapy.
were also obtained.
 Aruffo
 01-DEC-1992;
 US5830731-A.
 Misc-difference
 Misc-difference
 from a single patient undergoing extracorporeal interleukin-2 induction therapy. Clones pl35 (see V63456) and p90 (see X07372) were also obtained. A novel method for cloning cDNAs from mammal
 Example 10; Column 53-54; 75pp; English.
 (GEHO) GEN HOSPITAL CORP.
 immunotherapeutic applications,
 265
 1 EDGNVLKRSPEL 12
edgnvlkrspel 276
 l'Similarity
12; Conser
 including
 Conservative
 immune-mediated
 92US-0983647.
88US-0160416.
89US-0379076.
90US-0553759.
97US-0861205.
 97us-0861205
 /note=
 ~/note= "encoded by GGC"
159
 /note= "encoded
176
 /note= "encoded
171
 /note= "encoded 141
 /note= "encoded by GGG
 humans
 100.0%; Score 61; DB 19;
100.0%; Pred. No. 0.0016;
rive 0; Mismatches (
 "encoded
 infections,
 by GTC"
 bу
 þу
 by ccc"
 including the diagnosis and
 ATG"
 AAC"
 DB 19;
 diseases,
 Length 374;
 Indels
 and
 from mammalian
 disorders of
 0;
 Gaps
 0
```

RESULT

12

Χü

B43683 standard;

Protein;

399

Ā

B43683;

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RESULT
B43683
 Вb
 δÃ
 Y96226
ID Y96226 standard; Protein;
 Query Match
Best Local
 Matches
 receptor consist only of the extracellular portion of the receptor and are not glycosylated i.e. they do not have transmembrane domains or signal peptides. The recombinant proteins may be used in immunoassays to determine the immune status of patients with chronic diseases of the immune system, e.g. AIDS, systemic lupus erythematosus (SLE), multiple myeloma (MM), or rheumatoid arthritis. In addition, pharmaceutical compositions containing recombinant proteins may be used to treat or prevent autoimmune diseases, allergies or tumours, especially AIDS, prevent autoimmune diseases, allergies or tumours, especially AIDS,
 The present sequence is the human high affinity FC receptor, FCgammaRI. FCgammaRI is also known as CD64. FC receptors play an important role in defending the body against infections. First, pathogens are opsonised by serum immunoglobulins. The resulting complex then binds to cells expressing FC receptors. FCgammaRI molecules are expressed by monocytes and macrophages, but expression can also be induced on neutrophils and eosinophils. Upon FC receptor activation, immune effector pathways are activated, leading to immune response. The present sequence may be modified to produce recombinant versions. The recombinant FC
 eosinophil; HIV; IgG; immunosuppressive; antitheumatic; cytostatic; antiinflammatory; AIDS; SIE; multiple myeloma; rheumatoid arthritis systemic lupus ervihematosus.
 Y96226;
 Novel Fc receptor lacking transmembrane domains, a signal peptide, glycosylation, useful for diagnosing and treating immune disorders
 03-DEC-1998;
 systemic lupus
 Sequence
 Disclosure;
 WPI; 2000-367968/32
 (PLAC) MAX PLANCK GES
 03-DEC-1998;
 EP1006183-A1
 Homo sapiens
 Human high affinity Fc receptor, FcgammaRI.
 rheumatoid arthritis
 265
 13
 1 EDGNVLKRSPEL 12
 edgnylkrspel
 high affinity Fc receptor; FcgammaRI; immunoglobulin; ion; immune response; CD64; monocyte; macrophage; neut
 12;
 Similarity
 374 AA;
 Page 26-28; 60pp; English.
 Conservative
 (first entry)
 98EP-0122969
 98EP-0122969.
 276
 100.0%;
 or MM
 FOERDERUNG WISSENSCHAFTEN
 374
 0
 Score 61; DB 21;
Pred. No. 0.0016;
Mismatches 0
 DB 21;
 0;
 rheumatoid arthritis;
 Length 374;
 Indels
 0,
 Gaps
 0,
```

08-FEB-2001

(first entry)

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망
 Query Match
Best Local
 Matches
 polypeptides, antibodies, agonists and antagonists from the present invention may be used to treat immune disorders by activating or inhibiting the proliferation, differentiation or mobilisation of immune cells, to treat disorders of haematopoietic cells, autoimmune disorders, allergic reactions, graft versus host disease and organ rejection, modulate haemostatic or thrombolytic activity, modulate inflammation, cancers, cardiovascular disorders, neurological disease and bacterial or viral infections: The peptides, nucleotides, antibodies, agonists and antagonists may be also be used in drug screens. C78449 to C78457 and
 antithyroid; antiallergic; antibacterial; antiviral; dermatological; neuroprotective; cardiant; thrombolytic; coagulant; nootropic; vastixpoic; antipsoriatic and antianglogenic. The polynucleotides an polypeptides can be used for preventing, treating or ameliorating me conditions and diagnosing pathological conditions. Polynucleotides,
 C77607
B43398
 Sequence
 antagonists may be also B44240 represent sequence
 B43398 to B44239. The proteins can have activities based on and cells the genes are expressed in Example of activities cytostatic; proliferative; vulnerary; immunomodulator; antic
 Claim 11;
 Novel isolated nucleic acids comprising sequences useful for treating or diagnosing e.g. cancer-
 290
 invention.
 antiasthmatic; antirheumatic; antiarthritic; antiinflammatory;
 WPI; 2000-587533/55
N-PSDB; C77892.
 Rosen CA,
 (HUMA-) HUMAN GENOME SCI INC
 12-MAR-1999;
 08-MAR-2000;
 21-SEP-2000
 WO200055350-A1
 Homo sapiens
 neurological
 vasotropic; antipsoriatic; antiangiogenic;
immune disorder; haematopoietic cell disor
 dermatological;
 antidiabetic;
 diagnosis;
 Human; cancer associated
 Human cancer associated protein sequence SEQ ID NO:1128.
 allergic reaction;
 antiinflammatory;
 haemostatic;
 1 EDGNVLKRSPEL 12
edgnvlkrspel 301
 l Similarity
12; Conser
 to C78448 encode the human cancer associated proteins given in
 Page 1739-1740; 2352pp; English.
 -399 AA;
 Ruben SM;
 Conservative
 cytostatic;
 2000WO-US05882.
 thrombolytic; cardiovascular disorder; infection;
 antiasthmatic;
 99US-0124270.
 ntiasthmatic; antirheumatic; antiarthritic; antiviral; y; antithyroid; antiallergic; antibacterial; cardiant; neuroprotective; thrombolytic; coagulant; nouropoic; ipsoriatic; antiangiogenic; gene therapy; inflammation;
 sequences
 haematopoietic cell
 aematopoletic cell disorder; autoimmune disorder;
graft versus host disease; organ rejection;
 100.0%;
 drug screening.
 proliferative;
 gene; cancer
 0;
 Score 61; D
Pred. No. 0.
 Mismatches
 in drug screens.
in the exemplifi
 exemplification
 vulnerary;
 antigen;
 DB 21;
0.0017;
 The polynucleotides and
 detection;
 Length
 immunomodulator;
 encoding peptides
 of.
 antidiabetic;
 399;
 cancer;
 ٥,
 include:
 the tissues
 present
 Gaps
 medical
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X X X X E
 XI
 Misc-difference
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 Misc-difference
 Misc-difference
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 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Macrophage; FCRI; cell surface antigen; human; immunoselection; panning; immunodiagnosis; diagnosis; immunotherapy; gene therapy; immune disorder; infection; asthma; immune-complex disease; amyloidosis; multiple sclerosis; parasitic disease; Fc receptor.
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Homo sapiens
 19-DEC-2000
 Y96183 standard; Protein; 344.AA.
 macrophage-specific
 (first entry)
 /note=
 /note=
- 55
 /note=
220
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216
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268
 /note=
117
 /note=
136
 /note=
134
 /note=
129
 /note=
126
 /note=
123
 /note=
116
 139
 64
 .60
 Location/Qualifiers
'note= . "encoded by
 note=
 /note=
 note=
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 'note=
 "encoded by
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 "encoded by
 "encoded
 "encoded by
 "encoded by
 "encoded by
 "encoded by
 "encoded by
 "encoded by CAG"
 "encoded by TCC"
 "encoded
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 "encoded
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 "encoded by
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 "encoded
 FCRI
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 by GTG"
 γd
 by AAT"
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 Ьy
 by AAT
 Ьy
 bу
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 Ьy
 γď
 bу
 GGT
 AAC"
 TTC"
 AAT"
 CGT"
 CCA"
 GTT"
 cce"
 ACC"
 GAA"
 CAT"
 TTC"
 CAA"
 GCA'
 CCC
 CAC"
 cge"
 ACC"
 GCG"
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"encoded by CCT'

sapiens

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 Ş
 Query Match
Best Local S
Matches 11
 01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
23-MAR-1990;
13-JUL-1990;
 Macrophage; FCRI; cell surface antigen; human; immunoselect panning; immunodiagnosis; diagnosis; immunotherapy; gene th immune disorder; infection; asthma; immune-complex disease;
 the invention designed to isolate CSA nucleic acids. The method based upon transient expression of a CSA in eukaryotic cells and physical selection of cells expressing the antigen by adhesion to
 The present sequence is that of a human macrophage specific FcRI, as deduced from cDNA clone pl35 (see A50631), which was isolated from a cDNA library expressed in COS cells using a novel method of from a cDNA library expressed in COS cells using a novel method of
 Isolated nucleic acid molecule encoding the CD19 cell surface antigen, useful for immunodiagnosis and immunotherapy of immune-mediated infections or disorders, e.g. asthma, immune-complex disease, parasitic
 Human macrophage-specific FCRI.
 Y96134;
 Y96134 standard; Protein; 374 AA.
 surfaces other than macrophages, protein of FCRI and a receptor 1
 animals, including humans
 (panning on) an antibody-coated substrate such as a culture dish CSA nucleic acids isolated by the method of the invention, and the coated to the invention of the invention.
 WPI;
 Stamenkovic I,
 amyloidosis;
 19-DEC-2000 (first entry)
 Sequence
 ability to interfere with such bonding,
 sclerosis.
 Example 10; Column 53-55;
 diseases
 N-PSDB; A50631.
 (GEHO) GEN HOSPITAL CORP.
 28-OCT-1998;
 immune-complex disease, amyloidosis, parasitic diseases or multiple sclerosis. FCRI is a high affinity receptor for the Fc portion of IGG, normally located on the cell surfaces of macrophages. The ability to interfere with such bonding, or to cause it to occur on surfaces other than macrophages, is useful in therapy. A fusion protein of FCRI and a receptor ligand will be helpful to increase
 proteins they encode, are useful
 29-AUG-2000
 US6111093-A.
 265
 mmunotherapeutic applications, including the diagnosis and
 Local Similarity 91.
nes 11; Conservative
 1 EDGNVLKRSPEL 12
||| |||||||
| 65 edgmvlkrspel 276
 15
 potencies
 2000-586382/55.
 344
 multiple
 immune-mediated
 of antibodies
 92US-0983647.
88US-0160416.
89US-0379076.
90US-0498809.
90US-0553759.
 $
 98US-0181612
 Seed
 86.9%;
 В;
 sclerosis;
 75pp; English.
 These disorders
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 in therapy.
 Score 53;
Pred. No.
 infections,
 Mismatches
 parasitic disease;
 for immunodiagnostic and
 DB 21;
0.046;
 diseases, and disorders in ers include asthma,
 ۲
 immunoselection;
 Length 344;
 Indels
 Fc receptor
 gene therapy;
 0;
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 Gaps
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WPI; 2000-586382/55.
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01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
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13-JUL-1990;
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N-PSDB; A50592.

Isolated nucleic acid molecule encoding the CD19 cell surface antigen, useful for immunodiagnosis and immunotherapy of immune-mediated infections or disorders, e.g. asthma, immune-complex disease, parasitic diseases.

Example 10; Column 53-55; 75pp; English.

The present sequence is that of a human macrophage specific FCRI, as deduced from CDNA clone pl35 (see A50592), which was isolated from a CDNA library expressed in, COS cells using a novel method of the invention designed to isolate CSA nucleic acids: The method is CB based upon transient expression of a CSA in eukaryotic cells and CC physical selection of cells expressing the antigen by adhesion to CC (panning on) an antibody-coated substrate such as a culture dish. CC (coanning on) an antibody-coated substrate such as a culture dish. CC (coanning on) an applications, including the invention, and the CC proteins they encode, are useful for immunodiagnostic and coated infections, including the diagnostic and coated infections, including the diagnostis and coated infections, including the diagnostis and coated infections, include asthma, including humans. These disorders include asthma, coated on the cell surfaces of macrophages. The CC infection of the coated on the cell surfaces of macrophages. The CC surfaces other than macrophages, is useful in therapy. A fusion CC protein of FCRI and a receptor ligand will be helpful to increase xxx.

S XXCCCCCCCCCCCCCCCCXXXX P P P P XXD

Sequence 374 AA;

δÃ Query Match 86.9 Best Local Similarity 91.7 Matches 11; Conservative 86.9%; Score 53; DB 21; Length 374; Pred. No. 0.051; Indels 0; Mismatches 1; Indels 0; Gaps 0,

В 1 EDGNYLKRSPEL 12 ||| ||||||| || 265 edgmvlkrspel 276

Search completed: June Job time: 205 sec 4, 2001, 12:13:32

GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
 Minimum DB seq
Maximum DB seq
 Result
 Database
 Total number of hits satisfying chosen parameters:
 Searched:
 Scoring table:
 Sequence:
 Title:
Perfect score:
 Run on:
 OM protein -
 ŏ
 Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
 Score
 protein search, using sw model
 length: 0
length: 2000000000
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2: /cgn2_6/ptodata/:
3: /cgn2_6/ptodata/:
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Gapop 10.0
 US-09-284-107-27
61
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US-08-368-704C-37

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US-08-368-704C-36

US-08-368-704C-36

US-08-394-189B-5

PCT-US9-3-05701-20

PCT-US93-05705-5
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 Description
 Sequence
 Sequence
 Sequence
 Sequence
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US-08-332-562A-94
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Query Match

82.0%;

Score 50; DB 2;

Length 91;

linear

83.3%;

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 Matches
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Best Local
 GENERAL INFORMATION:
 Sequence 11, Approximately No. 5840683
 Best Local Similarity 83.3 Matches 10; Conservative
 GENERAL INFORMATION:
 TELEFAX: (303) 499-8089
NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
 HYPOTHETICAL: NO ORIGINAL SOURCE:
 MOLECULE TYPE:
 200 EDGSVLKR 207
 TELECOMMUNICATION INFORMATION: TELEPHONE: (303) 499-8080
 CURRENT APPLICATION DATA:
 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
 ATTORNEY/AGENT INFORMATION:
 CORRESPONDENCE ADDRESS:
 APPLICANT:
 ITLE OF INVENTION: Peptides Inhibiting the Oncogenic Action ITLE OF INVENTION: of P21 Ras
 TOPOLOGY: 11
 NAME: Ferber, Donna M.
REGISTRATION NUMBER: 33,
REFERENCE/DOCKET NUMBER:
 1 EDGNVLKR 8
 APPLICATION NUMBER: US/08/531,525 FILING DATE: 21-SEP-1995
 CLASSIFICATION:
 COUNTRY:
 LENGTH:
 ADDRESSEE:
 77 EDSSVLKRSPEL 88
 1 EDGNVLKRSPEL 12
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 Similarity
 amino acid
 80303
 Boulder
 Application US/08718270A
 Colorado
 Application US/08531525
 213 amino acids
Hlavka, Joseph J.
Pincus, Matthew R.
 E: Greenlee and Winner, P.C. 5370 Manhattan Circle, Suite
 Arabidopsis thaliana
 Abajian, Henry B. Kende, Andrew S.
 Hlavka, Joseph J. Pincus, Matthew R.
 linear
 No. 58406831e, John F.
 PatentIn Release #1.0, Version #1.30
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 21-SEP-1995
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Pred: No.
 Pred. No. 0.0086;
 Mismatches
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 36;
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 Indels
 Indels
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 Gaps
 Gaps
 В
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Query Match
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 US-08-737-825-8
 Patent No.
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 APPLICANT:
APPLICANT:
 TOPOLOGY: linear MOLECULE TYPE: pro HYPOTHETICAL: NO ORIGINAL SOURCE:
 TELEFAX: (303) 499-808
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SEQUENCE CHARACTERISTICS:
LENGTH: 213 amino acid
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 PRIOR APPLICATION DATA:
 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
 200 EDGSVLKR 207
 TELECOMMUNICATION INFORMATION:
 ATTORNEY/AGENT INFORMATION:
 PRIOR APPLICATION DATA:
 SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
 CORRESPONDENCE ADDRESS:
 APPLICANT: Kende, P
TITLE OF INVENTION:
TITLE OF INVENTION:
 APPLICANT: No. 59104781e, APPLICANT: Abajian, Henry
 STRANDEDNESS:
 TYPE: amino acid
 NAME: Ferber, Donna M.
REGISTRATION NUMBER: 33,878
REFERENCE/DOCKET NUMBER: 78
 APPLICATION NUMBER: US 60/004,091 FILING DATE: 21-SEP-1995
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FILING DATE: 21-SEP-1995
 FILING DATE: 20
CLASSIFICATION:
 1 EDGNVLKR 8
 ORGANISM:
 TELEPHONE:
 APPLICATION NUMBER:
 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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 ZIP: 80303
 COUNTRY:
 CITY: Boulder
 ADDRESSEE:
 INFORMATION:
 Colorado
 213 amino acids
 STEWART,
 5370 Manhattan
 BYCROFT, BARRIE WALSHAM
WILLIAMS, PAUL
 COX; ANTHONY RICHARD JOHN HOLDEN, MATTHEW THOMAS GEOFFREY PORTER, LAUREN ELIZABETH
 SEBAIHIA, MOHAMMED
 Conservative
 Arabidopsis thaliana
 linear
 Kende, Andrew
 (303) 499-8080
 Greenlee, Winner and Sullivan, 70 Manhattan Circle, Suite 201
 protein
 20-SEP-1996
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 499-8089
GORDON SIDNEY ANDERSON BIRNIE
GENES INVOLVED IN THE BIOSYNTHETIC PATHWAY OF CARBAPENEM
12
 GEORGE PEACOCK COPELAND
SIMON JAMES
 59.0%;
 Peptidomimetics Inhibiting the Oncogenic Action of P21 52
 US/08/718,270A
 John
 Score 36;
Pred. No.
 78-95
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 Ras
 Gaps
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ORRESPONDENCE ADDRESS

Birch,

Stewart, Kolasch & Birch,

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ORIGINAL SOURCE:
ORGANISM: Era
US-08-737-825-8
 US-08-394-189B-6
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Best Local Similarity 70.v
Conservative
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OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
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FILING DATE: 24-FEB-2005
 COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
 SEQUENCE CHARACTERISTICS:
LENGTH: 288 amino acids
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 TELECOMMUNICATION INFORMATION: TELEPHONE: (703) 205-8000 TELEPAX: (703) 205-8050
 SOFTWARE: MICROSOFT WORD97
CURRENT APPLICATION DATA:
 ATTORNEY/AGENT INFORMATION:
 137 DGKTLKRVPE 146
 ORRESPONDENCE ADDRESS
 PPLICANT: Shaham, Shai
ITLE OF INVENTION: REI
ITLE OF INVENTION: BET
 UMBER OF SEQUENCES:
 PLICANT:
 CITY: Boston
STATE: MA
 LENGTH:
 NAME: SVENSSON, LEONARD R. REGISTRATION NUMBER: 30,330
 COMPUTER:
 CLASSIFICATION:
 APPLICATION NUMBER: FILING DATE: 03-JUN
 COUNTRY:
 STREET:
 STRANDEDNESS:
 REFERENCE/DOCKET NUMBER:
 ADDRESSEE:
 2 DGNVLKRSPE 11
 E OF INVENTION:
 02110
 T: P.O. Box 747
Falls Church
 amino acid
 Application US/08394189B
 24834
 Virginia
 176 Federal Street
 USA
 Yuan, Junying
 Horvitz, Robert
 Erwinia carotovora
 IBM Compatible
 Clark & Elbing LLP
 linear
 03-JUN-1997
 protein
 GENE,
 RELATEDNESS OF HUMAN INTERLEUKIN-1
BETA CONVERTASE GENE TO A C. ELEGA
 59.0%;
70.0%;
 US/08/737,825
 INHIBITORY
 0
 1009-0105P
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 US-08-258-287B-37; Sequence 37, Ap
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 Matches
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 GENERAL INFORMATION:
 NFORMATION FOR SEQ ID NO:
 FILING DATE: 24-JUN-1993
ATTORNEY/AGENT INFORMÁTION:
NAME: BUGAÍSKY, Lawrence B.
REGISTRATION NUMBER: 35,086
REFERENCE/DOCKET NUMBER: 06
 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Competible
OPERATING SYSTEM: PC-DOS/MS-DOS
 TOPOLOGY: linear MOLECULE TYPE: protein
 SEQUENCE CHARACTERISTICS:
LENGTH: 479 amino acid
 APPLICATION NUMBER: 07/8
FILING DATE: 12-JUN-1992
ATTORNEY/AGENT INFORMATION:
 SOFTWARE: Patentin Rel
CURRENT APPLICATION DATA:
 TELECOMMUNICATION INFORMATION:
 PRIOR APPLICATION DATA:
 REFERENCE/DOCKET NUMBER: 01
RELECOMMUNICATION INFORMATION:
TELEPHONE: 617-428-0200
 ORRESPONDENCE ADDRESS:
 NUMBER OF SEQUENCES:
 ITLE OF INVENTION:
 TELEFAX:
 APPLICATION NUMBER: FILING DATE: 10-JUN CLASSIFICATION: 435
 TELEPHONE:
 APPLICATION NUMBER:
 CITY: Washington
 NAME: Bieker-Brady, Kristina REGISTRATION NUMBER: 39,109
 APPLICATION NUMBER: (FILING DATE: 20-NOV-J
 APPLICATION NUMBER: (FILING DATE: 12-JUL-)
 CLASSIFICATION:
 COUNTRY:
 STRANDEDNESS:
 ADDRESSEE:
 1 EDGNVLKRSPEL 12
 20005
 amino acid
 D.C.
 Application US/08258287B
 479 amino acids
 E: Sterne, Kessler, Goldstein & Fox 1100 New York Avenue, Suite 600
 617-428-7045
 Yuan, Junying
 USA
 Miura, Masayuki
 24-JUN-1993
 10-JUN-1994
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ID NO:
 Programmed Cell Death Genes and Proteins
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50.0%;
 Release #1.0, Version #1.25
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 08/282,211
 US 08/080,850
 US/08/258, 287B
 07/984,182
 01997/211001
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 Score 35;
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LENGTH:

amino acid

497 amino acids

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 US-08-258-287B-37
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Best Local Similarity
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LENGTH: 497 amino acids
TYPE: amino acid
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PRIOR APPLICATION DATA: APPLICATION NUMBER: US 08/080,850
FILING DATE: 24-UN-1993
ATTERNEY/AGENT INFORMATION:
 tent No.
 REGISTRATION NUMBER: 35,086
REFERENCE/DOCKET NUMBER: 0609.3920002
TELECOMMUNICATION INFORMATION:
465 QGANILKOMPEL 476
 SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
 APPLICANT: Yuan, Junying
APPLICANT: Miura, Masayuki
TITLE OF INVENTION: Programmed Cell Death Genes and Proteins
NUMBER OF SEQUENCES: 95
 PRIOR APPLICATION DATA:
 CORRESPONDENCE ADDRESS:
 465 QGANILKQMPEL 476
 TELEFAX: (202) 371
TELEX: 248636 SSK
 TELEPHONE:
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FILING DATE: 10-JUN-1994
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OPERATING SYSTEM: PC-DOS/MS-DOS
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 1 EDGNVLKRSPEL 12
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 DDRESSEE:
 [TY: Washington
 20005
 6087160
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 D.C
 Application US/08368704C
 E: Sterne, Kessler, Goldstein & Fox 1100 New York Avenue, Suite 600
 Conservative
 both
 USA
 Conservative
 both
 (202) 371-2600
 4-JAN-1995
N: 435
 371-2540
 57.4%;
50.0%;
 57.4%;
 US 08/258,287
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 Score 35; DI
Pred. No. 55;
 Score 35; DB
Pred. No. 55;
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 DB 3;
 Length 497;
 Length 497
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 Gaps
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RESULT

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 Patent No.
 GENERAL INFORMATION:
 Patent No.
 NEORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
 APPLICANT: Yuan, Junying APPLICANT: Miura, Masayuki TITLE OF INVENTION: Program
 GENERAL INFORMATION:
 CORRESPONDENCE ADDRESS:
 NUMBER OF SEQUENCES:
 STATE: D.C. COUNTRY: US ZIP: 20005
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PRIOR APPLICATION NUMBER: US 08/080,850
 SOFTWARE: Patentin Rel
 TELEPHONE: (202) 371-2600
 FILING DATE: 24-JUN-1993 ATTORNEY/AGENT INFORMATION:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
 STREET: 1100 New CITY: Washington
 CORRESPONDENCE ADDRESS
 APPLICANT: Yuan, Junying
APPLICANT: Miura, Masayuki
TITLE OF INVENTION: Programmed Cell Death Genes and Proteins
 Local Similarity
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 ADDRESSEE:
 1 EDGNVLKRSPEL 12
 NAME: Bugaisky, Lawrence B.
REGISTRATION NUMBER: 35,086
REFERENCE/DOCKET NUMBER: 0609.3920001
 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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 LENGTH:
 COUNTRY:
 STREET:
 ADDRESSEE:
 36,
 6, Application US/08368704C 6087160
 Washington
 6, Application US/08258287B 6083735
 amino acid
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 248636 SSK
 E: Sterne, Kessler, Goldstein & 1100 New York Avenue, Suite 600
 USA
 E: Sterne, Kessler, Goldstein & 1100 New York Avenue, Suite 600
 (202)
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 PatentIn Release #1.0, Version #1.25
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 10-JUN-1994
 371-2540
 Programmed Cell Death Genes and Proteins
 57.4%;
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 equence 5, Application US/08394189B atent No. 5962301
 TELEFAX: (202) 371-2540
TELEX: 248636 SSK
INFORMATION FOR SEQ ID NO: 36:
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LENGTH: 503 amino acid
 SOFTWARE: FastSEQ for Windows Version 2.0 CURRENT APPLICATION DATA:
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 ORRESPONDENCE ADDRESS
 471 QGSNILKQMPEL 482
 TELECOMMUNICATION INFORMATION
 FILING DATE: 24-JUN-1993
ATTORNEY/AGENT INFORMATION:
 CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
US 08/258,287
 PPLICANT:
 CURRENT APPLICATION DATA:
 FILING DATE: 20-NO APPLICATION NUMBER:
 STATE:
 NAME: Bugaisky, Lawrence B
REGISTRATION NUMBER: 35,080
 OPERATING SYSTEM:
 COMPUTER:
 COUNTRY:
 STREET:
 REGISTRATION NUMBER: 35,086
REFERENCE/DOCKET NUMBER: 0609.3920002
 APPLICATION NUMBER:
 APPLICATION NUMBER: US 0 FILING DATE: 10-JUN-1994
 APPLICATION NUMBER:
 ZIP: 02110
 ADDRESSEE:
 TLE OF INVENTION: BETA CONVERTASE GENE TO A C. ELEGANS CELL DEATH TLE OF INVENTION: GENE, INHIBITORY PORTIONS OF THESE GENES AND...
 TOPOLOGY:
 TELEPHONE:
 1 EDGNVLKRSPEL 12
 LE OF INVENTION:
 I: 503 amino acids amino acid
amino acid
GY: both
 Boston
 ΑM
 176 Federal Street
 USA
 Conservative
 Shaham, Shai
 Yuan, Junying
 Horvitz, Robert
 IBM Compatible
 Clark & Elbing LLP
 (202) 371-2600
 Diskette
JMBER: 07/897,788
12-JUN-1992
 4-JAN-1995
 57.4%;
 RELATEDNESS OF HUMAN INTERLEUKIN-1
BETA CONVERTASE GENE TO A C. ELEGAI
 Dos
 US/08/368,704C
 US 08/080,850
 3; Mismatches
 Score 35; DE
Pred. No. 56;
 DB 3;
 3; Indels
 Length 503;
 0;
 Gaps
 0,
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Query Match .
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 Query Match
Best Local Similarity
Watches 6; Conserve
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 PCT-US93-05701-20
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PCT-US93-05701-20
 Sequence 20, Applicat GENERAL INFORMATION:
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 TELECOMMUNICATION INFORMATION: TELEPHONE: 617-428-0200
 SEQUENCE CHARACTERISTICS:
 CORRESPONDENCE ADDRESS:
 TITLE OF INVENTION:
 APPLICANT:
 NUMBER OF SEQUENCES:
473 OGSNILKOMPEL 484
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REFERENCE/DOCKET NUMBER: 01997/211001
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OPERATING SYSTEM:
 COUNTRY:
 STRANDEDNESS:
 TYPE: amino acid TOPOLOGY: linear
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 1 EDGNVLKRSPEL 12
 ADDRESSEE:
 ENGTH:
 02139
 Cambridge
 Massachusetts
 Application PC/TUS9305701
 505 amino acids
 77 Massachusetts Avenue
 Massachusetts Institute of Technology
VENTION: Cloning and Characterization of Cell Death Genes
 Conservative
 ss: single
linear
 Conservative
 at every Xaa
 Massachusetts Institute of Technology
 57.4%;
 57.4%;
 20:
 Score 35; DB Pred. No. 56;
 Score 35; DB Pred. No. 56;
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 Length 505;
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 Indels
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 Gaps
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RESULT 12 PCT-US93-05705-5

Sequence 5, Application PC/TUS9305705 GENERAL INFORMATION:

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FILING DATE: 19930714
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LENGTH: 505 amino acids
TYPE: amino acids
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 GENERAL INFORMATION:
 REGISTRATION NUMBER: 33,367
REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-8850
 FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Warren M. Cheek, Jr.
NAME: Warren M. Cheek, Jr.
33,367
 SOFTWARE: Wordperfect 5.1
CURRENT APPLICATION DATA:
APPLICATION UNMEER: U5/08/588,985
FILING DATE: January 19, 1996
 STREET: BUJ
STREET: Washington
CITY: Washington
CTATE: D.C.
 CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
 COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette,
 NUMBER OF SEQUENCES: CORRESPONDENCE ADDRESS
 FEATURE:
 MOLECULE TYPE: protein
 COMPUTER READABLE FORM:
 COMPUTER: IBM COM
OPERATING SYSTEM:
 473 QGSNILKQMPEL 484
 CORRESPONDENCE ADDRESS:
 TITLE OF INVENTION:
 TLE OF INVENTION
 DDRESSEE:
 TOPOLOGY:
 1 EDGNVLKRSPEL 12
 MEDIUM TYPE:
 COUNTRY:
 STREET:
 ADDRESSEE:
 20005
 02139
 Cambridge
 Massachusetts
 E: Wenderoth, Lind & Ponack
805 Fifteenth Street, N.W.,
 E: Massachusetts Institute of Technology 77 Massachusetts Avenue
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 Massachusetts Institute of Technology ENTION: Inhibitors of Ced-3 and Related Proteins
 January
DN: 536
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 CDNA OF DOCK180 GENE AND DOCK180 PROTEIN
 PCT/US93/05705
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د:
 Score 35; DE
Pred. No. 56;
 Mismatches
 #700
 DB 5;
 3; Indels
 Length 505;
 0;
 Gaps
Query Match 57.4
Best Local Similarity 66.7
Matches 8; Conservative
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; FRAGMENT TYPE:
; ORIGINAL SOURCE:
US-08-971-988-2
 Query Match Best Local Similarity Conserve
 US-08-971-988-2
 ; FRAGMENT TYPE:
; ORIGINAL SOURCE:
US-08-588-985-2
 Patent No. 5786461
 INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 1865 amino acids
 GENERAL INFORMATION:
 ANTI-SENSE:
 HYPOTHETICAL:
 MOLECULE TYPE:
 TELEPHONE: 202-371-8850
 ATTORNEY/AGENT INFORMATION:
NAME: WAITEN M. Cheek, Jr.
REGISTRATION NUMBER: 3, 367
REFERENCE/DOCKET NUMBER:
 CURRENT APPLICATION DATA:
 LENGTH: 1000
TYPE: amino acid
 CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
 COMPUTER READABLE FORM: MEDIUM TYPE: Diskette,
 CORRESPONDENCE ADDRESS
 NUMBER OF SEQUENCES:
 APPLICANT: Michiyu
TITLE OF INVENTION:
 ANTI-SENSE:
 MOLECULE TYPE:
HYPOTHETICAL:
 169 EDGNIL -- DPEL 178
 STRANDEDNESS:
 TOPOLOGY:
 APPLICATION NUMBER: US/0 FILING DATE: 17-NOV-1997
 OPERATING SYSTEM:
 COMPUTER:
 COUNTRY:
 CITY: Washington
 STREET:
 STRANDEDNESS:
TOPOLOGY: lir
 ADDRESSEE
 1 EDGNVLKRSPEL 12
 ENGTH:
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 amino acid
 E: Wenderoth, Lind & Ponack
805 Fifteenth Street, N.W., #700
 1865 amino acids
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 Michiyuki MATSUDA et al.
'ENTION: CDNA OF DOCK180 GENE AND DOCK180 PROTEIN
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 IBM Compatible
 linear
 protein
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 MS-DOS
 57.4%;
 08/588,985
 US/08/971,988
 5.25 inch, 500 kb
 1; Mismatches
 Score 35; DB 1;
Pred. No. 2.5e+02
 Length 1865;
 Indels
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57.4%; Score 35; DB 1; 66.7%; Pred. No. 2.5e+02; tive 1; Mismatches 1

Length 1865;

Indels

2;

Gaps

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Page:
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HYPOTHETICAL: NO ORIGINAL SOURCE: ORIGINAL SOURCE: ORGANISM: Plasme US-08-568-459A-12
Search completed: June 4, 2001, 12:14:32 Job time: 225 sec
 밁
 Дb
 RESULT 15
US-08-568-459A-12
 Query Match
Best Local Similarity
7; Conserv
 REFERENCE/DOCKET NUMBER: NIH:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 235-8550
TELEFAX: (619) 235-0176
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
LENGTH: 2710 amino acids
 equence 12, Application US/08568459A
 ENERAL INFORMATION:
 2 DGNVLKRSPE 11
||||| ||:
1680 DGNVLSVSPK 1689
 ZIP: 92660
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
 NUMBER OF SEQUENCES: 3
 CURRENT APPLICATION DATA:
 CLASSIFICATION: 435
ITTORNEY/AGENT INFORMATION:
NAME: Israelsen, Ned
REGISTRATION NUMBER: 29,6
 PPLICANT: Su, Xin-zhaun
PPLICANT: Wellems, Thomas E.
ITLE OF INVENTION: BINDING DOMAINS FROM PLASMODIUM VIVAX
ITLE OF INVENTION: AND PLASMODIUM FALCIPARUM ERYTHROCYTE BINDING PROTEINS
 169 EDGNIL--DPEL 178
 ADDRESSEE: Knobbe Martens Olson & Bear
STREET: 620 Newport Center Drive 16th Floor
CITY: Newport Beach
STATE: California
 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
 COUNTRY: US
ZIP: 92660
 TYPE: amino acid
STRANDEDNESS: si
 APPLICATION NUMBER: FILING DATE: 07-DE
 1 EDGNVLKRSPĒL 12
 Sim, Kim L.
Chitnis, Chetan
Chiler, Louis H.
Miller, Louis B.
Peterson, David S.
 Plasmodium falciparum
 linear
 07-DEC-1995
N: 435
 single
 protein
 57.48;
70.08;
 US/08/568,459A
 29,655
 NIH121.001CP1
 Score 35; DB 2;
Pred. No. 3.9e+0;
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## ALIGNMENTS

C;Species: Homo sapiens (man)
C;Date: 03-Apr-1992 #sequence\_revision 03-Apr-1992 #text\_change 21-Jan-2000-C;Accession: A41357; S03019

FC gamma (IgG) receptor I (high affinity)
N;Alternate names: CD64

form b -

A;Title: Isolation and expression of functional high-affinity Fc receptor complementa A;Reference number: A41357; MUID:89100284

R; Allen, J.M.; Seed, 1 Science 243, 378-381,

4135,, : Seed, B. : 381, 1989

Accession: A41357

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FC gamma (IgG) receptor I-A (high affinity) precursor - human N;Alternate names: CD64 (Species: Homo sapiens (man) (C;Species: Homo sapiens (man) (C;Date: 30-Dec-1991 #sequence_revision 06-Sep-1996 #text_change 21-Jul-2000 (C;Accession: A39878; T10304; B41357; S03018; T57525 (R;van de Winkel, J.G.J.; Ernst, L.K.; Anderson, C.L.; Chiu, I.M. J. Biol. Chem. 266, 13449-13455, 1991 (A;Title: Gene organization of the human high affinity receptor for IgG, Fcgra, R;reference number: A39878; MUID:91302383
 RESULT
A39878
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 F;117-170/Domain: immunoglobulin homology <IMM>
 A;Cross-references: EMBL:X14355; NID:g31333; PIDN:CAA32536.1; PID:g31334
A;Note: the authors translated the codon AcT for residue 25 as Ala
C;Superfamily: Fr gamma receptor I; immunoglobulin homology
C;Keywords: alternative splicing; immunoglobulin receptor; transmembrane protein
 A; Molecule type: mRNA
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A;Accession: S03019
 R;Allen, J.M.; Seed, B.
Nucleic Acids Res. 16, 11824, 1988
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A;Cross-references: GB:X14355; GB:M21090; NID:g31333; PIDN:CAA32536.1; PID:g31334
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 1 EDGNVLKRSPEL 12
|||||||||
265 EDGNVLKRSPEL 276
 100.0%;
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 Score 61; DB 2; ... Pred. No. 0.00086;); Mismatches 0;
 Length 344;
 Indels
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A;Reference number: A46480; MUID:92166399
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 A; Residues:
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 д
 Qγ
 A46480
 c gamma (IgG) receptor high affinity mouse; Alternate names: high affinity IgG receptor Species: Mus musculus (house mouse)
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A;Molecule type: mRNA
A;Residues: 1-24,'S',26-337,'T',339-374 <AL2>
A;Cross-references: EMBL:X14356; NID:g31331; PIDN:CAA32537,1; PID:g31332
R;Perez, C.; Wietzerbin, J.; Benech, P.D.
Mol. Cell. Biol. 13, 2182-2192, 1993
A;Title: Two cis-DNA elements involved in myeloid-cell-specific expressions.
 Molecule
 Status:
 Osman,
 Query Match
Best Local S
Matches 12
 A; Title: Nucleotide sequence of three cDNAs A; Reference number: S03018; MUID:89098339
 Map position: 1q21-1q2;
Superfamily: Fc gamma
 Cross-references: GB:S57204; NID:g298692;
 ;Reference number: I57525; MUID:93204964;Accession: I57525
 Science 243, 378-381,
 ;59,78,152,159,163,195,240/Binding site: carbohydrate
 ;Gene: GDB:FCGR1A; CD64
 A;Cross-references: GB:M63830; R;Porges, A.J.; Redecha, P.B.;
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293-313/Domain: transmembrane #status predicted <TMM>
 Reference number: Accession: B41357
 Title: Isolation and expression of functions, Reference number: A41357; MUID:89100284
 Title: Novel Fc gamma
 cleic Acids Res.
 Residues:
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Residues: 1-24,'S','
 Status: translated from GB/EMBL/DDBJ
 Accession: S03018
 Allen, J.M.;
 Accession:
 lross-references: GB:X14356;
 Molecule type: mRNA
 Reference number:
 ross-references: GB:L03418; NID:g184840; PIDN:AAA36049.1; PID:g292169
 265
 N.; Kozak, C.A.; McKenzie, I.F.;
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e type: DNA
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 .M.; Seed, B.
ids Res. 16, 11824, 1988
 Similarity 100.
12; Conservative
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 Seed, B.
 1921-1921
 Redecha, P.B.;
. 90, 2102-2109,
 .26-337,'T',339-374 <AL1>
3B:X14356; GB:M21091; NID:931331; PIDN:CAA32537.1;
 1989
 receptor I; immunoglobulin homology
 100.0%;
 receptor I family 577; MUID:93055454
 involved in myeloid-cell-specific expression
 GB:M63835; NID:g180277; PIDN:AAA35678. Doebele, R.; Pan, L.C.; Salmon, J.E.;
 0;
 1992
 Score 61;
Pred. No.
 of functional high-affinity Fc receptor complementary
 Mismatches
 oprotein; immunoglobulin receptor; predicted <SIG>
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 gene products in
 0.00094;
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 for
 DB 1;
 the human high affinity Fc receptor (Fc
 mouse high
 Length 374;
 Indels
 (Asn)
 human mononuclear cells
 (covalent)
 affinity
 21-Jan-2000
 0;
 Gaps
 .1; PID:g180279
.Kimberly, R.P.
 FC
 #status predicte
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 transmembrane
 gamma
 and
 0
 gamma
 RH
 C; Accession: S77922
R; Dornelas, M.C.; Lecharny, submitted to the EMBL Data submitted to number: S77922
 A:Cross-references: EMBL:X99696; NID:g1480077; PIDN:CAA68027.1; PID:g1480078 C:Superfamily: kinase-related transforming protein; protein kinase homology
 shaggy-like protein kinase iota (EC 2.7.1.-)
C:Species: Arabidopsis thaliana (mouse-ear cr
C:Date: 23.Jul-1997 #sequence_revision 01-Auc
 A; Molecule type: mRNA
 RESULT
S77922
 A;Residues:
 A; Status: preliminary
 A; Accession: S77922
 В
 δÃ
 C; Keywords: traif; 37-53/Domain:
 A; Map position: 12R
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 A; Gene:
 A; Molecule type: DNA
A; Residues: 1-112 <DUZ>
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 A; Reference number:
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 Qy
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 Immunol.
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 Local
 1 EDGNVLKRSPEL
 1-407 <DOR>
 S55946
 Similarity
inase-related transforming protein;
phosphotransferase; protein kinase
 Lecharny, A.;
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Library,

August 1996

(mouse-ear cress

01-Aug-1997 #text\_change 20-Jun-2000

Arabidopsis thaliana

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probable membrane protein YLR390w - yeast (Saccharomyces N;Alternate names: hypothetical protein L8084.10 C;Species: Saccharomyces cerevisiae C;Date: 23-Aug-1995 #sequence_revision 19-Oct-1995 #text_C;Accession: S55946
 A;Cross-references: EMBL:U19729; NID:g625097; A;Experimental source: strain S288C (AB972)
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A;Description: The sequence of S. cerevisiae cosmid 8084.
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R; Sears,
 A;Title: Molecular cloning and expression A;Reference number: A43511; MUID:90111035
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 Local Similarity 66. nes 8; Conservative
 274 EDSSVLKRSPEL 285
 sequence extracted from D.W.; Osman, N.; Tate,
 transmembrane protein
ain: transmembrane #status predicted
 Osman, N.;
4, 371-378,
 Conservative
 sequence: S55944
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 12
 82.0%;
 Tate,
1990
 2;
 В.;
 NCBI backbone
 Score 42;
Pred. No.
 Score 50;
Pred. No.
 Mismatches
 McKenzie,
 Mismatches
 of
 PIDN:AAA40056.1;
lobulin homology
 DΒ
 PID:g625107;
 the mouse high affinity
 .13;
 (NCBIN:85205, NCBIN:85208, I.F.C.; Hogarth, P.M.
 <TMM>
 'n
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 Length 112;
 #text_change 06-Feb-1998
 Length 404;
 Indels
 Hogarth,
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 PID:g200753
0;
Gaps
 Gaps
 FC
 NCBIN: 85212,
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밁
 QV
 C; Superfamily: maltose permease
 밁
 R; Prins, J.B.; Todd, J.A.;
Science 260, 695-698, 1993
 Fc gamma (IgG) receptor high affinity - mouse
C;Species: Mus musculus (house mouse)
C;Date: 02-Jul-1996 #sequence_revision 02-Jul-1996 #text_change 21-Jan-2000
 뮹
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 Query Match
Best Local Similarity
Thes 8; Conserv
 C: Accession: I48471
 F; 68-329/Domain: F; 76-84/Region: F
 conserved hypothetical protein VC0302 [imported] - Vibrio cholerae (group Ol strain
 A; Molecule type: mRNA
A; Residues: 1-336 < RES>
 , R.R.; Mekalanos, J.J.;
lature 406, 477-483, 2000
 Residues: 1-396 <H
 ;Reference number: I48471; MUID:93242399;Accession: I48471
 ;Cross-references: GB:AE004119; GB:AE003852; NID:g9654718; PIDN:AAF93476.1; GSPDB:GN00;
;Experimental source: serogroup Ol; strain N16961; biotype El Tor
 ;Title: DNA Sequence of both chromosomes of the cholera pathogen Vibrio cholerae;Reference number: A82035; MUID:20406833;Accession: H82339
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Best Local Similarity
Matches 7; Conserv
 ;Gene: VC0302
 ;Status: preliminary; translated from GB/EMBL/DDBJ
 Title: Linkage on chromosome 3 of autoimmune diabetes and defective Fc receptor for
 Status: preliminary
 Heidelberg, J.F.; Eisen, J.A.; Nelson, W. nardson, D.; Ermolaeva, M.D.; Vamathevan,
 Date: 18-Aug-2000 #sequence_revision 20-Aug-2000 #text_change 15-Sep-2000
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 Matches
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 383 QDGNALKAEPE 393
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 1 EDGNVLKRSPE 11
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 Todd, J.A.; Rodrigues, N.R.;
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protein kinase ATP-binding mot
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 12
 290
 .; Venter, J.C.; Fraser,
 67.2%;
 68.9%;
72.7%;
 65.6%;
 <u>ب</u>
 Score 41;
Pred. No.
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 Score 42; DB
Pred. No. 4.2;
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 Mismatches
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 Ghosh,
 DB
5.2;
 Clayton, R.A.; Gwinn, M.L.; Dodson, R.J
Bass, S.; Qin, H.; Dragoi, I.; Sellers,
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 Length 336
 Length 396;
 Length 407;
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 P.M.; Wicker,
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A25233
replication protein -
C; Species: Bacillus su
C; Date: 05-Oct-1988 #s
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 A; Gene:
 C; Date:
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probable lo
C; Species:
 Genetics 148, 1021-1029, 1998
A;Title: Theme and variation among silencing proteins in
A;Reference number: Z16686; MUID:98198828
 A;Gene: APE1307
C;Superfamily: Synechocystis long-chain-fatty-acid--CoAC;Keywords: acid-thiol ligase; coenzyme AF;71-606/Domain: acetate--CoA ligase homology <ACL>
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6; Conserve
 A;Description: involved in both silencing C;Keywords: transcription regulation
 R;Astrom, S.U.; Rine, U.
R;Astrom, S.U.; Rine, U.
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 mating type silencing
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A;Experimental source: strain Kl
 A; Molecule type: DNA
A; Residues: 1-651 <KAW>
 A; Reference number: A; Accession: D72605
 A; Title: Complete genome
 R;Kawarabayasi, Y.; Hino, Y.; awa, H.; Takamiya, M.; Masuda, DNA Res. 6, 83-101, 1999
 R;Kawarabayasi,
 C;Date: 20-Aug-1
C;Accession: D72
 A;Status: preliminary
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Matches 7
 ;Species: Kluyveromyces marxianus var. lactis, Can;Date: 20-Sep-1999 #sequence_revision 20-Sep-1999
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 Function:
 432
 SIR4
 1 EDGNVLKRSPEL
 long-chain-fatty-acid--CoA ligase
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 EDGEILMRSPAM
 Similarity 7; Conser
 T09481
 Conservative
 Conservative
 Hino, Y.; Horikawa, H.; Y
M.; Masuda, S.; Funahashi,
 enome sequence of an aerobic hyper-thermophilic Crenarchaeon, Aero
A72450; MUID:99310339
 #sequence_revision
 regulator SIR4 -
 443
 12
 58.3%;
 62.3%;
54.5%;
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Pred. No. 84;
 Score
Pred.
 Mismatches
 Mismatches
 regulator
 20-Aug-1999 #text_change 03-Nov-2000
 No.
 yeast (Kluyveromyces marxianus var. lactis)
 actis, Candida sphaerica
 and telomere length maintenance
 16;
 Yamazaki,
11, T.; Tana
 (EC 6.2.1.3) APE1307 [similarity]
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 PID:g3002789
 Tanaka,
 #text_change 11-May-2000
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 Length 1314,
 Length 651
 S.; Haikawa, Y.;
aka, T.; Kudoh, Y.
 Indels
 ligase;
 Saccharomyces cerevisiae and
 acetate--CoA ligase
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 Gaps
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 .; Jin-no, K.;
Y.; Yamazaki,
 0
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EDGNVSEKTPQ 717

subtilis #sequence\_

Bacillus subtilis plasmid pIM13

revision 05-Oct-1988 #text\_change 10-Dec-1999

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C;Accession: A61069
R;Dyke, K.G.H.; Curnock, S.P.
FEMS Microbiol. Lett. 58, 209-216,
 C;Species: Staphylococcus aureus
C;Date: 31-Dec-1993 #sequence_revision
 A; Accession:
 replication protein REP - Staphylococcus aureus
 B,
 A; Reference number:
 A; Title: The nucleotide
 C; Superfamily:
 C; Accession: A
R; Monod, M.; I
J. Bacteriol.
 Ъ
 Qy
 ; Experimental source:
 N; Molecule type: DNA
 replication
 C; Superfamily: plasmid replication
 A;Title: Sequence and properties of piM13, A;Reference number: A91819; MUID:86250583
A;Accession: A25233
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 Title: Replication properties of pIM13, Reference number: A91847; MUID:88032835; Accession: A29827
 Experimental source: plasmid pE5 Catchpole, I.; Thomas, C.; Davie, Gen. Microbiol. 134, 697-709, 1
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 Reference number: A46568; MUID:89036120
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 Bacteriol.
 101
 101
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 Local
 The nucleotide sequence of Staphylococcus aureus
 EEGNIIKRKTGVLMLNPEL
 S.J.;
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 EEGNIIKRKTGVLMLNPEL 119
 plasmid
 Similarity
8; Conser
 protein REP - Staphylococcus aureus
 Similarity
8; Conser
 Denoya, C.; Dubnau, 167, 138-147, 1986
 plasmid
 Monod,
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 Conservative
 A61069
 5131-5139,
 replication protein
 plasmid
 sequence
 61.5%;
 represents
 61.5%;
 compared with conceptual translation
 Narayanan,
 12
 Davies,
 pT48
 of a
 4;
 1988; A.; Dyke,
 Score
Pred.
 Score 37.5; D
Pred. No. 9.9;
4; Mismatches
 D.
 the only long
 protein
 Mismatches
 small
 31-Dec-1993
 C.S.;
 37.5;
No. 1
 a naturally occurring
 REP
 cryptic
 REP
 Dubnau,
 macrolide-lincosamide-streptogramin
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 #text_change 10-Dec-1999
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 Length 158;
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 Length 146;
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RESULT
S19534
 C; Superfamily: lipocalin;
 Eur. J. Biochem. 202, 31-40, 1991
A;Title: Complete sequence and model for the C(1) subunit of
A;Reference number: S19534; MUID:92037648
 crustacyanin chain C1 - European lobster C; Species: Homarus gammarus (European lob C; Date: 19-Mar-1997 #sequence_revision 2:
 A; Molecule type: protein A; Residues: 1-181 < KEE>
 R;Keen,
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Best Local S
Matches 7
 C; Accession: S19534
 A; Molecule type: DNA
A; Residues: 1-162 < LAM>
 C; Superfamily:
 A; Reference number: A91821; A; Accession: B24497
 C; Genetics:
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B24497
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A;Molecule type: DNA
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C;Comment: This sequence represents
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Best Local S
Matches 8
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 A; Genome:
 C; Genetics:
 A; Title: Nucleotide sequence of
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 Accession:
 plication protein - Staphylococcus epidermidis plasmid pNE131
Species: Staphylococcus epidermidis
Date: 05-Jun-1988 #sequence_revision 05-Jun-1988 #text_change
 Matches
 Query Match
Best Local
 and pSN2.
 ampson, B.C.; Parisi, J.T.
Bacteriol. 167, 888-892, 1986
 101
 101 EEGNIIKRKTGALMLNPEL 119
2 DGNVLKRSPEL 12
 J.N.; Caceres,
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S
 1 EDGNVLKR---
 EEGNIIKRKTGVLMLNPEL 119
 EDGNVLKR----
 plasmid
 Similarity 7; Conserva
 Similarity 42.3
8; Conservative
 B24497
 Similarity 42.3
8; Conservative
 plasmid replication
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 #sequence_revision 25-Apr-1997
 I.; Eliopoulos, E.E.;
 ---SPEL 12
 replication
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 61.5%;
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 conceptual
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 the constitutive macrolide-lincosamide-streptogramin
 Score 37;
Pred. No.
 Score
 Score 37.5; I
Pred. No. 11;
4; Mismatches
 protein
 Score 37.5;
Pred. No. 1
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 protein
 the only long open
 Mismatches
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 DB
15;
 Zagalsky, P.F.; Findlay, J.B.C.
 ÐВ
 DB
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 2;
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 #text_change 10-Dec-1999
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 Indels
 Length 162;
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 Length 158;
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hypothetical protein F7H19.100 - Arabidopsis thaliana
C:Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 23-Apr-1999 #sequence_revision 23-Apr-1999 #text_change 23-Jul-1999
C;Accession: T05123
R;Bevran, M.; Peters, S.A.; van Staveren, M.; Dirkse, W.; Stiekema, W.; Bancroft, I.; Mew submitted to the Protein Sequence Database, July 1998
A;Reference number: 215399
A;Accession: T05123
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A;Residues: 1-268 (SEEV)
A;Croos-references: EMBL:AL031018
A;Experimental source: cultivar Columbia; BAC clone F7H19
C;Genetics:
A;Map position: 4
A;Introns; 47/3; 105/3; 161/3
A;Note: F7H19.100

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Matches 7; Conservative 1; Mismatches 3; Indels 0; Gaps 0;
Db 170 DGNLLNNYPEL 180

Search completed: June 4, 2001, 12:15:45
Job time: 278 sec
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|          |                |                                         | •          |                                      |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |                                       |                                                                | •                                     |               | \$ · ·  |
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|          | i i            |                                         |            |                                      | •                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              |                                       | •                                                              | · · · · · · · · · · · · · · · · · · · | ٠.            | •       |
|          | <del>i</del> · | • ()                                    |            |                                      |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |                                       |                                                                |                                       |               |         |
| •        |                |                                         |            |                                      | , , ;                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          |                                       |                                                                |                                       |               | x -30   |
|          |                |                                         | e .        | *                                    |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                | ٦                                     |                                                                |                                       | ,             | ( * -   |
|          |                |                                         | . 1        | *                                    |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                | •                                     |                                                                | •                                     |               |         |
| *        | •              |                                         | -          |                                      |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |                                       |                                                                | •                                     |               |         |
| *        |                |                                         | . '        |                                      |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                | ,                                     |                                                                |                                       |               |         |
| 31<br>2. |                | ,                                       |            |                                      |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |                                       |                                                                | * .                                   |               |         |
|          |                | 4                                       |            |                                      | 7                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              |                                       | • •                                                            | + 2                                   | T)            | * / L   |
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| F        | <del>-</del>   | San San San San San San San San San San |            | ·, ··· · · · · · · · · · · · · · · · |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |                                       |                                                                |                                       | * * . • . • . | *       |
|          | v č            |                                         |            |                                      |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |                                       |                                                                |                                       |               | *       |
|          |                | ••                                      |            | •                                    | •                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              | * .                                   | » ·                                                            |                                       |               | Ţ       |
| ·4.      |                | ,                                       |            |                                      |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                | 24                                    |                                                                | * .                                   |               |         |
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| - 6      |                |                                         |            |                                      |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |                                       |                                                                |                                       |               | ·       |
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| 4        |                | , af                                    | X ·        |                                      |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                | 9 8                                   |                                                                |                                       |               |         |
| K        |                | ¥                                       |            |                                      | * * *                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          |                                       |                                                                | - 1                                   |               | , · · • |
| 25       |                |                                         |            |                                      |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |                                       |                                                                |                                       |               | in .    |
| 7        |                |                                         |            |                                      |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |                                       |                                                                |                                       |               | *       |
| •        |                |                                         |            | . 8                                  | ·                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              | ** V                                  |                                                                |                                       |               |         |
| ·<br>·   |                |                                         |            |                                      | 3                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              |                                       |                                                                |                                       |               |         |
| r        |                | a *.                                    |            |                                      | ÷ 7                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            | *                                     |                                                                |                                       |               |         |
| - 1      |                |                                         | •          |                                      |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |                                       |                                                                | 440                                   |               |         |
| ris :    |                | `                                       | •          |                                      | . 4                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            |                                       |                                                                | •                                     |               |         |
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| er.      | * *            | 8.0                                     | ė          |                                      | 1. × 2. 2. 3. 3. 3. 3. 3. 3. 3. 3. 3. 3. 3. 3. 3.                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              |                                       | 4                                                              | * 4, * .                              |               |         |
|          |                |                                         |            | *                                    | •                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              |                                       |                                                                | * 1 81                                |               | •       |
| - 1      |                | **                                      |            |                                      | •                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              |                                       |                                                                | * *                                   |               |         |
|          |                | - CV =                                  |            | ^                                    | 1.0                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            |                                       |                                                                | . ***                                 | y (*)         |         |
| K.       | •              |                                         | *          | v V                                  | in the same                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    | +                                     |                                                                |                                       |               |         |
| 4        | -k-            |                                         |            |                                      |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |                                       |                                                                | 10.30                                 |               |         |
| 9        |                |                                         | -00-       |                                      |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |                                       |                                                                |                                       |               |         |
|          |                |                                         | A          |                                      |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |                                       |                                                                |                                       |               |         |
|          | 34             |                                         |            |                                      | The state of                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   | * 3. 13 ti                            |                                                                | 2. 11.16 ATT N. 2.                    | A 160 W       | £       |
|          | . *            |                                         |            |                                      |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |                                       | •                                                              |                                       | γ.            |         |
|          |                |                                         |            |                                      |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |                                       |                                                                |                                       |               |         |
|          |                | 1 - C                                   |            |                                      |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                | * *                                   |                                                                |                                       |               |         |
|          |                |                                         |            |                                      |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                | · · · · · · · · · · · · · · · · · · · | •                                                              |                                       |               | ,       |
| 3        |                |                                         |            |                                      |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |                                       |                                                                |                                       | . 47          |         |
| į.       |                |                                         |            |                                      |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |                                       | _ ^a/-                                                         |                                       | -             |         |
|          |                | . •                                     |            |                                      |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                | ·                                     |                                                                | •                                     |               |         |
| 7.       | :              |                                         | · •        | 4                                    | •                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              | × • •                                 | - <del>-</del> - <del>-</del>                                  | · · · · · · · · · · · · · · · · · · · | 3.            | 7       |
|          |                | *                                       |            |                                      | a de la companya de l | 1 · · · · ·                           | $e^{-1}(a_1^2,\ldots,a_{n-1}^2)=a_n^2(a_n^2,\ldots,a_{n-1}^2)$ | 4                                     |               |         |
| ą.       |                |                                         | 7. 1. just | 4                                    | ,                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              |                                       |                                                                |                                       |               | •       |
|          |                | Ž.                                      | N          |                                      | :6                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             |                                       |                                                                |                                       | (2)<br>(2)    |         |
|          |                | •                                       | 74         |                                      | · · · · · · · · · · · · · · · · · · ·                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          |                                       |                                                                |                                       |               | ٠. ز    |
|          | •              |                                         | 2,         |                                      |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |                                       |                                                                |                                       | ,             |         |
|          |                |                                         |            |                                      |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                | y.                                    | -14-                                                           |                                       | ,             | 52.     |
|          |                | `.                                      | 20         | •                                    | 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          | . 1                                   | , 1 7 × 1                                                      |                                       |               | rank.   |

```
Result
 Database
 Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
 Minimum DB
Maximum DB
 Total number of hits satisfying chosen parameters:
 Searched:
 Scoring table:
 Sequence:
 Title:
Perfect score:
 Run on:
 OM protein -
 õ
 118
119
120
120
120
120
130
130
130
130
130
 Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
 Score
 seg length: 0
seg length: 2000000000
 protein search, using sw
 Query
Match
 June 4, 2001, 12:24:07; Search time 37.56 Seconds (without alignments)
10.944 Million cell updates/sec
 BIOSUM62
Gapop 10.0 , Gapext 0.5
 SwissProt_39:*
 93435 seqs, 34255486 residues
 US-09-284-107-27
 1 EDGNYLKRSPEL 12
 GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd
 Length DB
 156
299
299
UBL1_HUMAN
UBL1_RAT
LAH1_SCHPO
 MRP_STRSU
YBAZ_ECOLI
HS11_CHERU
VP19_CRV
YE18_METJA
 ACTZ_PNECA
MTTF_HUMAN
IMPB_SALTY
PURA_VIBPA
Y4NK_RHISN
TOP1_XENLA
CHB_VIBHA
 RECF_STAAU
OPPC_MYCPN
 GFAP_BOVIN
HS11_ARATH
HS12_ARATH
LAXC_LACLA
LAXP_LACLA
 REMA_STAEP
CRC1_HOMGA
 FCG1_HUMAN
FCG1_MOUSE
KSGI_ARATH
 SSX3_HUMAN
ARA2_ARATH
 SUCC_ECOLI
HS12_ORYSA
 Ħ
 REMA_BACSU
 SUMMARIES
 model
 93435
 P41512
P13670
 Description
 rhizobium s
xenopus lae
vibrio harv
 streptococc
escherichia
 chenopodium
 vibrio para
 escherichia
 homo
 schizosacch
 homo sapien
 methanococc
 cymbidium r
 salmonella
 homo sapien
 pneumocysti
 mycoplasma
 staphylococ
 arabidopsis
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|   | 4     | 4     | 4        | 4          | 4     | 4     | ω     | ω     | ω     | ω            | ω          | ω          |  |
|---|-------|-------|----------|------------|-------|-------|-------|-------|-------|--------------|------------|------------|--|
|   | 5     | 4     | ω        | 2          |       | 0     | 9     | 8     | 7     | 6            | 5          | 4          |  |
|   | ·ω    | ω     | ω        | ω          | ω     | ω     | ω     | ų     | ω     | w            | ω          | :<br>س     |  |
|   | ω     | ω     | ω        | ω          | ω     | ω     | -     | -     | 4     | 4            |            | 4          |  |
|   | 54.1  | 54.1  | 54.1     | 54.1       | 54.1  | 54.1  | 55.7  | 55.7  | 55.7  | 55.7         | 55.7       | 55.7       |  |
| • |       |       |          |            |       |       |       |       |       |              |            |            |  |
|   | 291   | 252   | 215      | 189        | 181   | 151   | 581   | 985   | 922   | 798          | 569        | 565        |  |
|   | _     | _     | <b>-</b> | Н          | ب     | ب     | Н     | Н     | _     | <u>ن</u> ــر | <u>-</u>   | مبو        |  |
| : | YF92_ | YGHR_ | SC4_(    | VP11_RDVF  | VP11_ | HS11_ | VGLP_ | NAH1  | YKFO_ | L100_        | KIRI_      | KIRI       |  |
|   | SYNY. | _ECOI | CTDC     | _RDVE      | RDV2  | WHE!  | BEV   | YEAS  | _YEAS | _ADEC        | _SAL1      | ECOI       |  |
|   | 3     | H     | Ö        | .,         | _     | Ä     |       | ij    | ï     | ×            | Y          | Н          |  |
| - |       |       |          |            |       |       |       |       |       |              |            |            |  |
|   |       |       |          |            |       |       |       |       |       |              |            |            |  |
| _ |       |       |          |            |       |       |       |       |       |              |            |            |  |
|   | P729  | Q468  | P270     | 085442     | P288  | P128  | P230  | Q992  | P357  | P36856       | P061       | P082       |  |
|   | 70    | 42    | 12       | 42         | 90    | 10    | 52    | 71    | 36    |              |            |            |  |
| • | syne  | esch  | octo     | rice dwarf | rice  | trit  | bern  | saccl | sacci | avian adeno  | salmonella | escherichi |  |
|   | choc  | ericl | enc o    | dwa        | dwai  | icum  | e vi  | naron | naron | ade          | nel]       | erich      |  |
|   | /st   | 11a   | lof      | Œ          | îf    | ae    | su.   | ус    | īУC   | one          | ຜ          | lia        |  |
|   |       |       |          |            |       |       |       |       |       |              |            |            |  |

## ALIGNMENTS

RESULT 1
FCG1\_HUMAN RESULT

```
RESULT
FCG1_M
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 Query Match
Best Local S
 Matches
 1_MOUSE
FCG1_MOUSE
P26151;
01-MAY-1992
01-MAY-1992
15-JUL-1999
 SEQUENCE FROM N.A.

MEDLINE-92166399; PubMed-1531670;

MEDLINE-92166399; PubMed-1531670;

Osman N., Kozak C.A., McKenzie I.F., Hogarth P.M.;

"Structure and mapping of the gene encoding mouse gamma RI and chromosomal location of the human Fc.

J. Immunol. 148:1570-1575(1992).
 MEDLINE=90111035; PubMed=2136886; Sears D.W., Osman N., Tate B., Mc "Molecular cloning and expression receptor for IgG.";
 01-MAY-1992 (Rel. 22, Created)
01-MAY-1992 (Rel. 22, Last sequence
15-JUL-1999 (Rel. 38, Last annotati
HIGH AFFINITY IMMUNOGLOBULIN GAMMA
 CONFLICT
SEQUENCE
 DOMAIN
DOMAIN
DISULFID
DISULFID
 Eukaryota; Metazoa;
Mammalia; Eutheria;
 CARBOHYD
CARBOHYD
 CARBOHYD
CARBOHYD
 receptor for IgG
 SEQUENCE FROM N.A.
 Mus musculus
 FCGR1
 NCBI_TaxID=10090
 VARIANT
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 CARBOHYD
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 CARBOHYL
 DISULFID
 DOMAIN
DOMAIN
 VARSPLIC
 TRANSMEM
 DOMAIN
 SIGNAL
 265
 IgG-binding pro
Immunoglobulin
 Immunol.
 FUNCTION: BIRDS TO THE FC REGION OF IMMUNOGLOI AFFINITY RECEPTOR.
SUBCELLULAR LOCATION: TYPE I MEMBRANE PROTEIN TISSUE SPECIFICITY: MACROPHAGE-SPECIFIC.
 1 EDGNVLKRSPEL
 (FCRI)
 EDGNVLKRSPEL
 OR FCG1
 ch 100.0%; l Similarity 100.0%; 12; Conservative (
 (IGG
 144:371-378(1990).
 374
 (Mouse)
 240
333
 59
78
152
159
163
195
 43
124
212
 protein;
 25
 26
 ΑA;
 CONTAINS 3 IMMUNOGLOBULIN
 STANDARD;
 FC RECEPTOR I).
 domain;
 276
 12
 Chordata;
Rodentia;
 92
175
267
85
168
 25
42605
 59
 26
 Last sequence up
Last annotation
LOBULIN GAMMA FC
 Receptor; Transmembrane; Glycoprotein; Signal; Alternative splicing; Polymórphism.
 expression
 , WM
 IMMUNOGLOBULIN-LIKE C2-TYPE
 0;
 N-LINKED
N-LINKED
 Craniata; Veri
Sciurognathi;
 /FTId=VAR_003954.
S -> T (IN CAA32536).
2C2AA8103ECF16E6 CRC64;
 BY SIMILA
 CYTOPLASMIC (POTENTIAL).
IG-LIKE C2-TYPE DOMAIN.
IG-LIKE C2-TYPE DOMAIN.
IG-LIKE C2-TYPE DOMAIN.
 Score 61;
Pred. No.
 HEKKYTSSLQEDRHLEEELKCQEQKEEQLQEGVHRKEPQGA
 N-LINKED
 N-LINKED
 N-LINKED
 PRT;
 /FTId=VAR_003953
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 splicing; Polymorphism
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thi; Muridae;
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 IMMUNOGLOBULIN GAMMA FC
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 I PRECURSOR (FC-GAMMA
 Length 374;
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 Euteleostomi;
Murinae; Mus
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 Query Match
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 Q39012;
01-OCT-2000
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 _ARATH
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 STRAIN=CV. COLUMBIA; TISSUE=Root;
Dornelas M.C., Schwebel-Dugue N., Thomas
"Three new cDNAs related to SGC/GSR-3 (SK
kinase-3) from Arabidopsis thaliana.";
 Arabidopsis thaliana (Mouse-ear cress).
Eukaryota; Viridiplantae; Embryophyta;
Magnoliophyta; eudicotyledons; core eud
Brassicales; Brassicaceae; Arabidopsis.
 01-OCT-2000 (Rel. 40, I
01-OCT-2000 (Rel. 40, I
SHAGGY-RELATED PROTEIN
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 ASK9 OR GSK1 OR T2D23.9
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 PIR; A43511; A43511.
PIR; A46480; A46480.
HSSP; P12319; 1ALT.
 274
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 This SWISS-PROT entry is copyright between the Swiss Institute of Bioi
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 -!- CAUTION:
 gG-binding
 1 EDGNVLKRSPEL
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 PF00047; ig; 3. inding protein; Receptor;
 Similarity 83.
10; Conservative
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Gene Register
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 AAA40056.1;
 STANDARD;
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N-LINKED (GLCNAC...
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N-LINKED (GLCNAC...
 1;
 Score 50;
Pred. No.
 CYTOPLASMIC (POTENTIAL)
IG-LIKE C2-TYPE DOMAIN
IG-LIKE C2-TYPE DOMAIN.
IG-LIKE C2-TYPE DOMAIN.
 HIGH AFFINITY IMMUNOGLOBU RECEPTOR I.
EXTRACELLULAR (POTENTIAL)
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 7-LINKED (GLCNAC. .
1-LINKED (GLCNAC. .
1-LINKED (GLCNAC. .
1C4F0033842767E7 C
 Transmembrane; Glycoprotein; Signal;
 Mismatches
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 (EC 2.7.1.-) (ASK-IOTA).
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 407
 Nas M., Lecharny A., Kreis (SHAGGY/glycogen synthase
 Tracheophyta;
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(AUG-1997)

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 Phosphorylation; I
DOMAIN 70
NP_BIND 76
BINDING 99
ACT_SITE 195
SEQUENCE 407 AA
 Toriumi M., Yu G., Chan G., Chao Q., Com. L., Buehler E., Chao Q., Com. L., Altafi H., Brooks S., Buehler E., Chao Q., Com. Lam B., M Altafi H., Johnson-Hopson C., Khan S., Kim C., Lam B., M Hansen N.F., Johnson-Hopson C., Southwick A., Davis R.W., Balm C.J., Shinn P., Southwick A., Davis R.W.,
 EMBL;
 EMBL;
 Submitted (JUL-2000) to the EMBL/GenBank/DDBJ databases.
-!- FUNCTION: MAY MEDIATE EXTRACELLULAR SIGNALS TO REGULAT TRANSCRIPTION IN DIFFERENTIATING CELLS (BY SIMILARITY)
Monod M., Denoya C., Dubnau D.; "Sequence and properties of pIM13, a macrolide-lincosamide-streptogramin
 BACSU
 use
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 1
 Bacillus/Staphylococcus
NCBI_TaxID=1423;
 Pfam; .PF00069; pkinase; 1.
PROSITE; PS00107; PROTEIN_KINASE_ATP;
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 This SWISS-PROT entry is copyright. It is produced through a collaboration
 Medine=86250583; Pu. Monod M., Denoya C.,
 SEQUENCE FROM N.A.
 Bacteria; Firmicutes;
 Plasmid pIM13
 Bacillus subtilis.
 REPLICATION
 01-JAN-1990
15-DEC-1998
 01-JAN-1990
 P13968;
 REMA_BACSU
 InterPro
 InterPro; IPR000719;
 Mendel; 13957; Arath;2319;13957
 SEQUENCE, FROM N.A.
 21
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 SIMILARITY: BELONGS TO THE SER/THR FAMILY OF PROTEIN KINASES. CDC2/CDKX SUBFAMILY; GSK-3 SUBSUBFAMILY.
 (BY SIMILARITY)
 DGDALKRRPEL
 P24941; 1AQ1
 X99696; CAA68027.1; -.
 AC068143;
 Similarity 72.
8; Conservative
 PS00108; PROTEIN_KINASE_ST; PS50011; PROTEIN_KINASE_DOM;
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 AUTOPHOSPHORYLATED MAINLY ON THREONINE AND SERINE RESIDUES
 IPR002290; -.
 (Rel.
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 AND MAINTENANCE PROTEIN
 Serine/threonine-protein kinase;
 STANDARD;
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AAF82167.1;
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 Multigene family
354 PROTI
 PubMed=3087948;
 195
 46024 MW;
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 68.9%;
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 Created)
 Bacillus/Clostridium group;
 Etgu P., Lee J.M., Lenz C., Pham P., Sakai
n A., Chung M., Goldsmith A., Liu A., Smitl
Buehler E., Chao Q., Conn L., Conway A.B.,
ppson C., Khan S., Kim C., Lam B., Miranda
 group; Bacillus.
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1; Mismatches
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96BC4C53754A764C CRC64;
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 This SWISS-PROT entry is copyright. It is produbetween the Swiss Institute of Bioinformatics the European Bioinformatics Institute. There a
 Bacillus subtilis.";
J. Bacteriol. 167:138-147(1986)
 Staphylococcus aureus, and Staphylococcus simulans Plasmid pE5, and Plasmid pPV142.
Bacteria; Firmicutes; Bacillus/Clostridium group;
 EMBL; M13761; AAA98135.1; PIR; A25233; A25233.
 Somkuti G.A., Solaiman D.K.Y., Steinberg D.H.; Submitted (OCT-1997) to the EMBL/GenBank/DDBJ databases.
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 01-JAN-1990
15-DEC-1998
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 EMBL; M17990; AAA98227.1; -.
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 This SWISS-PROT entry is copyright. It is produced through a collaboration
 SPECIES-S
 SEQUENCE FROM N.A.
 "Replication properties of pIM13, a naturally occurring plasmid found in Bacillus subtilis, and of its close relative pE5, a plasmid native
 SPECIES=S.aureus;
MEDLINE=88032835;
 SEQUENCE FROM N.A.
 Bacillus/Staphylococcus group;
 REPL OR REP
 REPLICATION
 P13969;
 REMA_STAAU
 SEQUENCE
 DNA replication;
 EMBL;
 between
 Projan S.J., Monod M., Narayanan C.S.,
 NCBI_TaxID=1280,
 101 EEGNIIKRKTGVLMLNPEL 119
 Staphylococcus aureus.";
Bacteriol. 169:5131-5139(1987).
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 .simulans; STRAIN=13044; PLASMID=pPV142;
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 P80029;
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Ol-OCT-1994 (Rel. 30, Last annotation
CRUSTACYANIN C1 SUBUNIT
 Homarus gammarus (European lobster) (Homarus vulgaris).
Eukaryota, Metazoa; Arthropoda; Crustacea; Malacostraca;
Eumalacostraca; Eucarida; Decapoda; Pleocyemata; Astació
 _HOMGA
 "Complete sequence carotenoprotein, cr
 Findlay J.B.C.;
 Nephropoidea; Nephropidae; Homarus NCBI_TaxID=6707;
 CRC1
 STAEP
 MEDLINE=92037648; PubMed=1935978; Keen J.N., Caceres I., Eliopoulos
 SEQUENCE
 Homarus gammarus (European lobster)
 entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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 Lampson B.C., Parisi J.T.;
"Nucleotide sequence of the constitutive macrolide-lincosamide-streptogramin B resistance plasmid pNEI31 from Staphylococcus epidermidis and homologies with Staphylococcus aureus plasmids
 Bacteria; Firmicutes; & Bacillus/Staphylococcus
 REPL
 01-JAN-1988
15-DEC-1998
 P06735; Q52044;
01-JAN-1988 (Re
 EMBL; M12730; AAA98295.1;
 modified
 J. Bacteriol. 167:888-892(1986)
 SEQUENCE FROM N.A.
MEDLINE=86304180; PubMed=3091582;
 Plasmid pNE131
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 REPLICATION
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 Firmicutes; Bacillus/Clostridium grostaphylococcus group; Staphylococcus
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 P07460;
01-APR-1988
01-APR-1988
01-OCT-2000
 Oshima T., Aiba H., Baba T., Fujita K., Hayashi K., Honjo A., Ikemoto K., Inada T., Itoh T., Kajihara M., Kanai K., Kashimoto K., Kimura S., Kitagawa M., Makino K., Wasuda S., Miki T., Mizobuchi K., Mori H., Motomura K., Nakamura Y., Nashimoto H., Nishio Y., Saito N., Sampei G., Seki Y., Tagami H., Takemoto K., Wada C., Yamamoto Y., Yano M., Horiuchi T.;
 _ECOLI
 Pigment;
DISULFID
 MEDLINE-97426617; PubMed-9278503;
Blatther F.R., Plunkett G. III, Bloch C.A.,
Riley M., Collado-Vides J., Glasner J.D., R
Gregor J., Davis N.W., Kirkpatrick H.A., Go
 Yano M., Horiu
"A 718-kb DNA
 Mau B., Shao Y.; "The complete genome sequence of Escherichia coli K-12."; Science 277:1453-1474(1997).
 SEQUENCE FROM N.A.
MEDLINE=86104124; PubMed-3002435;
Buck D. Spencer M.E., Guest J.R.,
"Primary structure of the succiny."
 InterPro; IPR003057; -.
Pfam; PF00061; Lipocalin; 1.
PRINTS; PR01273; INTERTCOLOR.
PROSITE; PS00213; LIPOCALIN; FALSE_NEG.
 MEDLINE=97061202; PubMed=8905232;
 SEQUENCE FROM N.A.
 SEQUENCE FROM N.A. STRAIN-K12 / MG1655;
 coli."
 NCBI_TaxID=562;
 Bacteria; P
Escherichia
 Escherichia coli
 SUCCINYL-COA
 SEQUENCE
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 InterPro;
 COMPLEXES OF 16 SUBUNITS (ALPHA-CRUSTACYANIN).
TYPES OF SUBUNITS: A1, A2, A3, C1 AND C2.
-i- SUBCELLULAR LOCATION: EXTRACELLULAR, CARAPACE.
 COLORATION TO THE CARAPACE OF THE LOBSTER.
-!- SUBUNIT: OLIGOMER; CAN FORM DIMERS (BETA-CRUSTACYANIN);
 astaxanthin
 beta-crustacyanin,
 Biochemistry 24:6245-6252(1985)
 -! - SIMILARITY:
 -!- FUNCTION: BINDS
 Local Similarity
 73
 2 DGNVLKRSPEL
 _ECOLI
 DGNLLKRNGKL
 S19534; S19534.
 J. Biochem.
 Lipocalin;
 IPR000566; -
 Proteobacteria; gamma subdivision; Enterobacteriaceae;
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181 AA;
 (Rel. 07, Created)
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A SYNTHETASE BETA CHAIN (EC 6.2.1.5)
 Conservative
sequence of the Escherichia coli K-12 genome to the 12.7-28.0 min region on the linkage m
 STANDARD;
 BELONGS TO THE LIPOCALIN FAMILY.
 83
 202:31-40(1991).
IDS THE CAROTENOID
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 Transport.
 20667 MW;
 60.7%;
 the succinyl-CoA
 Score 37;
Pred. No.
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EMBL; J01619; AP
EMBL; AE000176;
EMBL; D90711; BP
 "ADP-binding site of Escherichia coli succinyl-CoA revealed by X-ray crystallography."; Biochemistry 39:17-25(2000)
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 MEDLINE-99141407; PubMed-9917402; Fraser M.E., James M.N., Bridger W.A., "A detailed structural description of
 coli at 2.5-A resolution.";
J. Biol. Chem. 269:10883-10890(1994).
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 Wolodko W.T., Fraser M.E., James M.N.G., Bridger W "The crystal structure of succinyl-CoA synthetase
 x-ray Crystallography (2:5 angstroms)
medline=94193797; PubMed=8144675;
 or send an
 MEDLINE=20092606; PubMed=10625475;
 synthetase.
 Electrophoresis 18:1259-1313(1997).
 STRAIN=K12 / EMG2;
MEDLINE=97443975; PubMed=9298646;
 SEQUENCE
SWISS-2DPAGE; P07460; COLI.
 "Comparing the predicted and observed properties
in the genome of Escherichia coli K-12.";
 Link A.J.,
 -RAY CRYSTALLOGRAPHY (3.3 ANGSTROMS).
 -RAY CRYSTALLOGRAPHY (2.3 ANGSTROMS)
 SUBUNIT: TETRAMER OF TWO ALPHA AND TWO BETA CHAINS.
MISCELLANEOUS: SUCCINYL-COA SYNTHETASE (SCS) OF E.COLI CATALYZES
ITS REACTION VIA THREE STEPS THAT INVOLVE PHOSPHORYL ENZYME AND
 ENZYME REGULATION: EXHIBITS TWO INTERESTING PROPERTIES: "SUBSTRAY SYNERGISM", IN WHICH THE ENZYME IS MOST ACTIVE FOR THE CATALYSIS OF ITS PARTIAL REACTIONS ONLY WHEN ALL THE SUBSTRATE SINDING SITUARE OCCUPIED, AND "CATALYTIC COOPERATIVITY" BETWEEN ALTERNATING ACTIVE SITES IN THE TETRAMER, WHEREBY THE INTERACTION OF SUBSTRATES (PARTICULARLY ATP) AT ONE SITE IS NEEDED TO PROMOTE
 MISCELLANEOUS: THE BETA-SUBUNIT CONTAINS THE ATTACHMENT SITES F
SUCCINATE. THE COMPLETE ACTIVE SITE IS PROBABLY LOCATED IN THE
REGION OF ALPHA- BETA CONTACT.
SIMILARITY: TO OTHER BETA SUBUNITS OF SUCCINYL-COA SYNTHETASES.
 SUBSTRATE-LEVEL PHOSPHORYLATION. IT CAN ALSO FUNCTION IN THE OTHER DIRECTION FOR ANABOLIC PURPOSES, AND THIS MAY BE PARTICULARLY IMPORTANT FOR PROVIDING SUCCINYL-COA DURING ANAEROBIC
 ENZYME-BOUND SUCCINYL PHOSPHATE AS INTERMEDIATES.
MISCELLANEOUS: DURING AEROBIC METABOLLSM IT FUNCTIONS IN THE
CITRIC ACID CYCLE, COUPLING THE HYDROLYSIS OF SUCCINYL-COA TO
SYNTHESIS OF ATP & THUS REPRESENTS AN IMPORTANT SITE OF
 European Bioinformatics Institute.
 GROWTH WHEN THE OXIDATIVE ROUTE
 CATALYSIS AT THE OTHER.
 ORTHOPHOSPHATE.
 CATALYTIC ACTIVITY:
 A24090;
 J01619; AAA23899.1;
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 3:137-155(1996)
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 Robison K.,
 20-APR-95.
02-AUG-99.
 0-JAN-00
 SYECSB
 285:1633-1653(1999)
 BAA35394.1;
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 Church G.M.;
 SUCCINATE + COA +
 FROM
 Escherichia coli succinyl-CoA
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 Wolodko W.T.;
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RESULT 10 SSX3\_HUMAN ID SSX3\_H

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 Matches
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 Query
 Nishi R., Hashimoto H., Uchimiya H., Kato A.;
Submitted (JUL-1992) to the EMBL/GenBank/DDBJ
-!- SUBUNIT: FORMS OLIGOMERIC STRUCTURES.
-!- SUBCELLUIAR LOCATION: CYTOPLASMIC.
 Oryza.
NCBI_TaxID=4530;
 ORYSA
 Pfam; PF00549; ligase-COA; l.
PROSITE; PS01217; SUCCINYL_COA_LIG_3; l.
Ligase; Tricarboxylic acid cycle; 3D-structure.
SEQUENCE 388 AA; 41392 MW; 09C429EC97A823CF CRC64;
 ECO2DBASE; E039.8;
EcoGene; EG10981; s
 Pfam; PF00011; HSP20; 1.
PROSTTE; PS01031; HSP20; 1.
Heat shock; Multigene family.
SEQUENCE 154 AA; 17376 MW;
 Oryza sativa (Rice).
Eukaryota; Viridiplantae; E
Magnoliophyta; Liliopsida;
 HS12_ORYSA
P31673;
 EMBL; D12635; BAA02160.1; PIR; JS0710; JS0710.
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 This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EN
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 InterPro; IPR000303;
 InterPro; IPR002068;
 SEQUENCE FROM N.A.
 17.4 KDA CLASS
 219
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 2 DGNVLKRSPEL
 SIMILARITY: PLANTS HAVE FOUR CLASSES OF HSP20: CLASS I AND I WHICH ARE CYTOPLASMIC; CLASS III WHICH IS IN THE CHLOROPLAST AND CLASS IV WHICH IS IN THE ENDOMEMBRANE. THIS PROTEIN BELO
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I HEAT SHOCK PROTEIN.
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 Magnoliophyta; eudicotyledo
Brassicales; Brassicaceae;
 CONFLICT
SEQUENCE
 Eukaryota; Viridiplantae;
 Arabidopsis thaliana
 ARA-2
 ARATH
 Q99909; O60223;
15-DEC-1998 (Rel
15-DEC-1998 (Rel
01-OCT-2000 (Rel
 PROSTIE; PS50806; KRAB_RELATED; 1.
Multigene family; Transcription regulation.
 This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its
 118
 or send
 MEDLINE=98021352; PubMed=937
Gure A.O., Tureci O., Sahin
Knuth A., Pfreundschuh M., C
 DOMAIN
 entities
 "SSX: a multigene family with several members transcribed
 testis and human cancer.
 SEQUENCE FROM
 sarcomas.";
 de Leeuw
 MEDLINE=96302330;
 Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
 human X chromosome
 Mammalia; Eutheria;
NCBI_TaxID=9606;
 SSX3 PROTEIN.
 SEQUENCE FROM N.A
 J. CARCET 72:965-971(1997).
FUNCTION: COULD ACT AS A MODULATOR OF TRANSCRIPTION.
SIMILARITY: BELONGS TO THE SSX FAMILY.
SIMILARITY: CONTAINS 1 KRAB-RELATED DOMAIN.
 EEGNVSKEVPE 128
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U90840; AAC05819.1;
TE; PS50806; KRAB_REI
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 b., Balemans M., Geurts van Kessel A.,
Kruppel-associated box containing the SSX gene (SSX3)
chromosome is not implicated in t(X;18) positive synov;
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O., Sahin U., Tsang S., schuh M., Old L.J., Chen
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PubMed=8697803;
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 Q28115;
01-NOV-1997
 SEQUENCE OF 197-227 FROM N.A.

SEQUENCE OF 197-227 FROM N.A.

Kirkpatrick B.W., Hart G.L., Moore S.S., Barendse W.;

Submitted (OCT-1993) to the EMBL/GenBank/DDBJ databases.

'- FUNCTION: GRAP, A CLASS-III INTERNEDIATE FILMARNIT, IS A CELL-
SPECIFIC MARKER THAT, DURING THE DEVELOPMENT OF THE CENTRAL
NERVOUS SYSTEM, DISTINGUISHES ASTROCYTES FROM OTHER GLIAL CELLS.
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 Eukaryota; Metazoa;
Mammalia; Eutheria;
 TISSUE=Subcommissural organ;
Bouchard P.;
 NCBI_TaxID=9913;
 Bovidae; Bovinae;
 Bos taurus (Bovine).
 GLIAL FIBRILLARY
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01-NOV-1997
 SEQUENCE FROM N.A.
 GTP-binding;
 EMBL; D01024; BAA00829.1;
PIR; J00907; J00907.
PIR; JS0639; JS0639.
HSSP; P28746; 1A2K.
 use by non-profit institumodified and this statement
 "Isolation and analysis of
of Arabidopsis thaliana.",
Gene 108:259-264(1991).
 InterPro; IPR001806; -
InterPro; IPR002380; -
 This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EN
 STRAIN=VARIOUS STRAINS; TISSUE=Leaf, MEDLINE=92084144; PubMed=1748311; Anai T., Hasegawa K., Watanabe Y., U
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 1 EDGNVLKR
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 Takahashi T., Komeda Y.; "Characterization of two genes encoding small heat-shock proteins
 Arabidopsis thaliana (Mouse-ear cress).
Arabidopsis thaliana (Mouse-ear cress).
Enkarvota; Viridiplantae; Embryophyta; Tracheophyta; Spermatophyta;
II;
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Pfam; PF00038; filament; 1.
 EMBL; Y08255; CAA69422.1; EMBL; L19867; AAA51413.1;
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 Arabidopsis thaliana.";
 STRAIN=CV. COLUMBIA;
MEDLINE=90158498; PubMed=2482931;
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 SIMILARITY: PLANTS HAVE FOUR CLASSES OF HSP20: CLASS I AND II WHICH ARE CYTOPLASMIC; CLASS III WHICH IS IN THE CHLOROPLAST AND CLASS IV WHICH IS IN THE ENDOMEMBRANE. THIS PROTEIN BELOW
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 Brassicaceae; Arabidopsis.
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 . 219:365-372(1989).
LOCATION: CYTOPLASMIC.
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01-FEB-1994 (Rel. 28, Last annotation
17.6 KDA CLASS I HEAT SHOCK PROTEIN (H
 HSP17.6.
Arabidopsis thaliana (Mouse-ear cress).
Arabidopsis thaliana (Mouse-ear cress).
Tracheophyta; Spermatophyta;
II;
 STRAIN-CV. COLUMBIA; TISSUE-Leaf;
MEDLINE-90016890; PubMed-2798141;
Helm K.W., Vierling E.;
"An Arabidopsis thaliana cDNA clone encoding
 Eukaryota; Viridiplantae; Embryophyta; Magnoliophyta; eudicotyledons; core euc Brassicales; Brassicaceae; Arabidopsis
 HS12_ARATH P13853;
 PROSITE; PS01031; HSP20; 1. Heat shock; Multigene family. SEQUENCE 156 AA; 17454 MW;
Heat shock;
SEQUENCE
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 PIR; S06074;
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PIR; JQ0351; JQ0351.
 EMBL; X16076; CAA34208.1;
 -1 - SIMILARITY: PLANTS HAVE FOUR CLASSES OF HSP20: CLASS I AND
 -!- SIMILARITY: BELONGS
 heat shock protein."
 NCBI_TaxID=3702;
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 Acids Res. 17:7995-7995(1989).
 Similarity
 IPR002068;
: Multigene family.
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CL Lactococcus lactis (Subsp. lactis)
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RN Huang D.C., Novel M., Huang X.F., Novel G.;
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Query Match
Best Local Similarity 41.7%; Pred. No. 26;
Matches 5; Conservative 5; Mismatches 2; Indels 0;

Qy 1 EDGNVLKRSPEL 12
Db 269 EEGNILEDKPQV 280

Search completed: June 4, 2001, 12:24:08

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GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.

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Post-processing: Minimum Match 0%
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Q91qe8 arabidopsis
 Q53734 staphylococ
 Q9nf41 drosophila
 Q06011 saccharomyc
 Q9kv55 vibrio chol
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 Q92495
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|-------------------|--------------------|--------------------|--------------------|--------|--------|--------------------|--------|--------------------|--------|--------------------|--------|--------------------|--------------------|--------|--------------------|--------------------|--------------------|--------|--------------------|--------------------|--------------------|--------|--------|-------------------|--------------------|
| 36                | 36                 | 36                 | 36                 | 36     | 36     | 36                 | 36     | 36                 | 36     | 36                 | 36     | 36                 | 36                 | 36     | 36                 | 36                 | 36                 | 36     | 36                 | 36                 | 37                 | 37     | 37     | 37                | 37                 |
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| <br>Q9VXP2        | Q9LQE3             | Q9N5D4             | Q9PAH0             | Q9XB56 | Q9SMF2 | P93440             | Q40867 | Q9ZP84             | Q9ZS24 | P93439             | Q9LDG9 | Q9M408             | Q9M4 09            | Q9M410 | Q9M411             | Q9M416             | Q9M417             | Q9M419 | 09M414             | Q9M412             | 016103             | Q9V8K9 | Q9V6K1 | 076251            | 017088             |
|                   |                    |                    |                    |        |        |                    |        |                    |        |                    |        |                    |                    |        |                    |                    |                    |        |                    |                    |                    |        |        |                   |                    |
|                   |                    |                    |                    |        |        |                    |        |                    |        |                    |        |                    |                    |        |                    | •                  |                    | •      |                    |                    |                    |        |        |                   |                    |
| Q9vxp2 drosophila | Q91qe3 arabidopsis | Q9n5d4 caenorhabdi | Q9pah0 xylella fas |        | N      | P93440 oryza sativ |        | Q9zp84 quercus sub |        | P93439 oryza sativ |        | Q9m408 quercus sub | Q9m409 quercus sub | _      | Q9m411 quercus sub | Q9m416 quercus sub | Q9m417 quercus sub |        | Q9m414 quercus sub | Q9m412 quercus sub | '016103 drosophila | w      | •      | 076251 drosophila | 017088 caenorhabdi |

## ALIGNMENTS

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 ID
 Q92495
 RESULT
 Query Match 100.0%; Score 61; DB 4; Best Local Similarity 100.0%; Pred. No. 0.0031; Matches 12; Conservative 0; Mismatches 0
Q92495
 "Novel Fc gamma receptor I tamily gecells.";
J. Clin. Invest. 90:2102-2109(1992).
 Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
 01-FEB-1997 (TrEMBLrel.
01-FEB-1997 (TrEMBLrel.
01-OCT-2000 (TrEMBLrel.
 Q92663
 MEDLINE=93055454; Put Porges A.J., Redecha Kimberly R.P.;
 INTERPRO: IPR003006; -. PFAM; PF00047; ig; 3. PF00047; ig; 3. PRODOM; PD002534; -; 1. SEQUENCE 374 AA; 426
 EMBL; L03418; AAA36049.1;
HSSP; P12319; IALT.
 Q92663;
 SEQUENCE FROM N.A.
 FC GAMMA RECEPTOR I.
 TISSUE=BLOOD;
 NCBI_TaxID=9606;
 265 EDGNVLKRSPEL 276
 1 EDGNVLKRSPEL 12
 N
 Fc gamma receptor I family gene products in human mononuclear
 (TrEMBLrel. 02, (TrEMBLrel. 02, (TrEMBLrel. 15,
PRELIMINARY;
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tha P.B., Doebele R.,
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42632 MW; D33D59398CEEA699 CRC64;
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PRT;
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375 AA
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 Salmon J.E.,
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RESULT
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 Yan Y., Li X., Wang A., Zhan
Molecular cloning and iden
high affinity Fc receptor for
Yet Immunol Immunopathol
EMBL; AF162866; AAF80477.1;
 Q9MZT0;
Q9MZT0;
01-OCT-2000
 092495;
01-FEB-1997 (TrEMBLrel. 02
01-FEB-1997 (TrEMBLrel. 02
01-OCT-2000 (TrEMBLrel. 15
FC GAMMA RECEPTOR TYPE I ()
Receptor
 Yan Y., Li X., Wan
 Bos taurus (Bovine).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutel
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora;
 Bovidae;
 NCBI_TaxID=9913
 01-OCT-2000 (TrEMBLrel.
01-OCT-2000 (TrEMBLrel.
 Structure.
 EMBL;
 SEQUENCE
 SEQUENCE OF 1-224 FROM MEDLINE=93018827; Pubme
 EMBL; M91555; AAA58414
 gamma receptor gene
 Benech P.
 Submitted (DEC-1992) to the
 Ezekowitz A.B.
 Benech P.D., Sastry K.N., Iyer R.R.,
 Eukaryota; Metazoa;
Mammalia; Eutheria;
 266
 *Definition of interferon gamma-response elements in a novel human
 TISSUE=BLOOD;
 SEQUENCE FROM N.A.
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M91554; AAA58414.1;
S45709; AAD13842.1;
S45707; AAD13842.1;
S45708; AAD13842.1;
S45708; AAD13842.1;
S45708; AAD13842.1;
 EDGNVLKRSPEL
 OR FC<GAMMA>RIB.
 PF00047;
 м91550;
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 Eutheria; Primates;
 (TrEMBLrel.
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AAA58414.1;
 176:1115-1123(1992)
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 g A., Zhang G.;
and identification of full-length
eceptor for bovine IgG (Fc gamma RI
 PubMed=1402657;
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 Borkova D., Minana B., "Sequencing the distal Submitted (FEB-1999) to
 Ballew R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M., Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S., Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P., Burtis K.C., Busam D.A., Butler H., Cadleu E., Center A., Chandra I., Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P., de Pablos B., Delcher A., Deng Z., Mays A.D., Dew I., Dietz S.M., Dodson K., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P. Durbin K.J., Evangelista C.C., Ferzac C., Ferziera S., Fleischmann W.
 SEQUENCE FROM N.A.
 01-MAY-1999
01-MAY-1999
01-MAY-2000
 George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N., Sutton G.G., Wortman J.F., Yandell M.D., Zhang Q., Chen L.X., Brandon R.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfeiffer B.D., Wan K.H., Doyle C., Baxter E.G., Helt G., Nelson C.R., Miklos G.L.G.
 Adams M.D., Celniker S.E., Holt R.A., E. Amanatides P.G., Scherer S.E., Li P.W.,
 MEDLINE=20196006; PubMed=10731132;
 SEQUENCE FROM N.A. STRAIN-BERKELEY;
 Pterygota; Neoptera; Ephydroidea; Drosoph
 Eukaryota;
 097172;
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 Drosophila
 FCP3C OR EG:140G11.3
 SEQUENCE
 266 DGNLIKRSPEL 276
 2 DGNVLKRSPEL
 J.F., Agbayani A., An H.-J., Andrews-Pfannkoch C.,
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 melanogaster
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Metazoa; Arthropoda;
 349 AA;
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 Drosophilidae;
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 S Y.-H.C., Blazej R.G., Champe M., Pfeiffer Baxter E.G., Helt G., Nelson C.R., Miklos
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 Mismatches
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 Evans C.A.,
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A., Galle R.
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Baldwin D.,
 Muscomorpha;
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Benos P.; Submitted ( EMBL; AE003

SEQUENCE

FROM N.A.

Submitted (FEB-1999) to the E EMBL; AE003426; AAF45849.1; -- EMBL; AL035395; CAB37612.1; -- FLYBASE; FBBG0000644; Fcp3C. SEQUENCE 99 AA; 11150 MW;

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OBF00050D1F11758 CRC64;

EMBL/GenBank/DDBJ databases.

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1 EDGNVLKRSPE : | | | | : | :

Saccharomyces cerevisiae (Baker's yeast).
Eukaryota; Fungi; Ascomycota; Saccharomycetes;
Saccharomycetaceae; Saccharomyces.

Saccharomycetales

SEQUENCE FROM N.A NCBI\_TaxID=4932;

PubMed=9169871;

01-NOV-1996

Q06011; Q06011

PRELIMINARY;

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CHROMOSOME XII COSMID

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STRAIN=S288C (AB972);
 Scharfe M., Scherens B., Scholler P., Schwager C., Schwarz Scholler P., Schwager C., Schwarz Sunderwood A.P., Urrestarazu L.A., Vandenbol M., Verhasselt F. Vierendeels F., Voet M., Volckaert G., Voss H., Wambutt R., Vierendeels F., Voet M., Volckaert A., Hani J., Wambutt R., Wedler H., Zimmermann F.K., Zollner A., Hani J., Handler B., Wedler H., Zimmermann F.K., Zollner A., Hani J., Wedler H., Wedler H., Zimmermann F.K., Zollner A., Hani J., Wedler H., Zimmermann F.K., Zollner A., Hani J., Wedler H., Wedler H., Zimmermann F.K., Zollner A., Wedler H., Zimmermann F.K., Zollner A., Wedler H., W
 Muller-Auer S., Nentwick v., Muller-Auer S., Purnelle B., Rechmann S.,
 STRAIN=S288C (AB972);
MEDLINE=97313267; PubMed-
Johnston M., Hillier L.,
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 Cherry J.M.;
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 Benes V. Bruckner M., Delius H., Dubois E., Dusterhoft A.,
Entian K.D., Floeth M., Goffean A., Hebling U., Heumann K.,
Heuss-Neltzel D., Hilbert H., Hilger F., Kleine K., Kotter P.,
Louis E.J., Messenguy F., Mewes H.W., Miosga T., Mostl D.,
 Waterston
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 Submitted (JAN-1995)
 Nature
 "The nucleotide sequence of Saccharomyces
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 ted (SEP-1997) to the EMBL/GenBank/DDBJ databases U19729; AAB82349.1; -. NCE 112 AA; 12515 MW; B469042AC2025AA7 CRC64;
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 Piravandi E.,
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 SEQUENCE FROM N.A.
STRAIN-CV. COLUMBIA;
Liu S.X., Vaysberg M., Etgu P.
 01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-CT-2000 (TrEMBLrel. 15, Last sequence update)
01-GCT-2000 (TrEMBLrel. 15, Last annotation update)
SHAGGY-LIKE PROTEIN KINASE IOTA (TZDZ3.9 PROTEIN)
ASK IOTA OR GSK1 OR TZDZ3.9.
ARABIOOPSIS thaliana (Mouse-bar Cress)
 Toriumi M., Yu G., Ch
Altafi H., Brooks S.,
 Eukaryota; Viridiplantae; Embryophyta; Trache
Magnoliophyta; eudicotyledons; core eudicots;
Brassicales; Brassicaceae; Arabidopsis.
 039012;
01-NOV-1996
 Submitted (OCT-1999) to the EMBL/GenBank/DDBJ databases EMBL; AL121805; CAB58072.1; SEQUENCE 299 AA; 32007 MW; C7224AFCDC0809B6 CRC64;
 Q9NF41
 Hansen N.F., Johnson-Hopson C., Nguyen M., Palm C.J., Shinn P.,
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TISSUE=ROOTS;
DOTTRELAS M.C., Schwebel-Dugue N., Thomas M., Lecharny A., Kreis
Plant Physiol. 113:306-306(1997).
 Q3901'2
 Piao H.L., Jang H.J.,
 STRAIN=COLUMBIA;
 "Sequencing
 SEQUENCE FROM N.A.
 Ephydroidea;
 Pterygota; Neoptera;
 Eukaryota;
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 01-0CT-2000
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 01-OCT-2000
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 Submitted
 NCBI_TaxID=7227;
 236
 82
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 Similarity 6; Conserv
 Cadieu, Dreano, Lelaure, Galibert F.; ing the distal X chromosome of Drosophila melanogaster."; d (OCT-1999) to the EMBL/GenBank/DDBJ databases.
 (AUG-1997)
 Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
1; Drosophilidae; Drosophila
 melanogaster (Fruit fly)
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Khan S., Kim C., Lam B., Mirana.
 Lim J.H.,
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 EMBL; AE004118; ATIGR; VC0302; -. SEQUENCE 396 A
 01-NOV-1999 (TrEMBLrel.
01-NOV-1999 (TrEMBLrel.
01-OCT-2000 (TrEMBLrel.
651AA LONG HYPOTHETICAL
 SEQUENCE FROM N.A.

STRAIN-EL TOR N16961 / SEROTYPE O1;

MEDLINE-20406833; PubMed-10952301;

Heidelberg J.F., Eisen J.A., Nelson W.C., Clayton R.A., Gwi
Dodson R.J., Haft D.H., Hackey E.K., Peterson J.D., Umayam

Gill S.R., Nelson K.E., Reickey E.K., Qin H., Richardson

Ermolaeva M.D., Vamathevan J., Bass S., Qin H., Dragoi I.,

McDonald L., Utterback T., Fleishmann R.D., Nierman W.C., W

Salzberg S.L., Smith H.O., Colwell R.R., MeKalanos J.J., Ve
 Q9YCF0
 Q9YCF0
 383
 Nature 406:477-483(2000)
 cholerae
 01-OCT-2000
01-OCT-2000
 PFAM; PF00069; pkinase; 1.
PROSITE; PS00107; PROTEIN KINASE_ATP; 1.
PROSITE; PS00108; PROTEIN_KINASE_DOM; 1.
PROSITE; PS50011; PROTEIN_KINASE_DOM; 1.
SEQUENCE 407 AA; 46024 MW; 96BC4C537
 HYPOTHETICAL
 Q9KV55;
 Q9KV55
 EMBL; AF019927; AAB71545.1;
EMBL; AC068143; AAF82167.1;
HSSP; P24941; 1JSU.
 'DNA sequence of both chromosomes
 NCBI_TaxID=666;
 Bacteria; Proteobacteria;
 Vibrio cholerae
 "The sequence of BAC T2D23 from Arabidopsis thaliana chromosome 1.", Submitted (JUL-2000) to the EMBL/GenBank/DDBJ databases.
 MENDEL; 13957; Arath; 2319; 13957.
 EMBL; X99696; CAA68027
 INTERPRO;
 INTERPRO;
 1 EDGNVLKRSPE 11 : | | | | | |
 21
 9
 2 DGNVLKRSPEL 12
 QDGNALKAEPE 393
 DGDALKRRPEL 31
 Similarity 63.77; Conservative
 8; Conserv
 IPR002290; -
 IPR000719
 PRELIMINARY;
 (TremBLrel. 15, Created)
(TremBLrel. 15, Last sequence update)
(TremBLrel. 15, Last annotation update)
 PRELIMINARY;
 Conservative
 PROTEIN VC0302
 AAF93476.1;
 43560
 65.6%;
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12, Last sequence update)
15, Last annotation update)
LONG-CHAIN-FATTY-ACID--COA
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 Score 42; DB
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 4D840465DBDC2757 CRC64
 96BC4C53754A764C CRC64;
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 Q9ZAS6;
01-MAY-1999
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 SEQUENCE
 PFAM; PF00496; SBP_bac_5; Lipoprotein.
 Mol. Microbiol. 28:1323-1334(1998).
EMBL; U78968; AAC67217.1; -.
INTERPRO; IPR000914; -.
 Podbielski A., Leonard B.A., "The group A streptococcal duptake of essential amino ac
 protease."
 MEDLINE=98343814; PubMed=9680220;
 STRAIN=CS10
 Streptococcus
 crenarchaeon, Aeropyrum pernix Kl.";
DNA Res. 6:83-101(1999).
 SEQUENCE FROM N.A.
 NCBI_TaxID=1314;
 Streptococcus
 Bacteria; Firmicutes;
 DPPA.
 SURFACE LIPOPROTEIN DPPA.
 Q9ZAS6
 SEQUENCE
 INTERPRO; IPR000873; PROSITE; PS00455; AM
 EMBL; AP000061; BAA80298.1; HSSP; P08659; 1LCI.
 Yamazaki J., Kushida N., Gyuchi A., Aoki K.,
Nomura N., Sako Y., Kikuchi H.;
"Complete genome ~~~~~
 Kawarabayasi Y., Hino Y., Horikawa H., Yamazaki
Jin-no K., Takahashi M., Sekine M., Baba S., An
Hosoyama A., Fukui S., Nagai Y., Nishijina K.,
Takamiya M., Masuda S., Funahashi T., Tanaka T.
 432
 MEDLINE-99310339; PubMed-10382966;
Kawarabayasi Y., Hino Y., Horikawa
 STRAIN-K1
 SEQUENCE FROM
 Aeropyrum.
NCBI_TaxID=56636;
 Aeropyrum pernix
Archaea; Crenarc
 APE1307.
 10
 1 EDGNVLKRSPEL
 EDGEILMRSPAM
 Similarity
9; Conserv
 Similarity 58. 7; Conservative
 p A streptococcal dipeptide essential amino acids and a
 Crenarchaeota;
 542 AA;
 651 AA;
 (TrEMBLrel. 10, (TrEMBLrel. 13,
 (TrEMBLrel.
 PRELIMINARY;
 pyogenes
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 12
 AMP_BINDING;
 61145 MW;
 63.1%;
56.2%;
 74662 MW;
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 Bacillus/Clostridium
 3.68;
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 Desulfurococcales;
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 Score 38.5;
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Pred.
 9CB9466D84F71FB6 CRC64;
 D641C4B159CD6754 CRC64;
 UNKNOWN_1
 Mismatches
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 01-OCT-2000
01-OCT-2000
01-OCT-2000
 Submitted (MAR-2000) to the EMBL/GenBank/DDBJ databases EMBL; AF247646; AAF66982.1; -. SEQUENCE 709 AA; 81941 MW; F1080D806D982ABF CRC64;
 Zea mays (Maize).
Eukaryota; Viridiplantae; E
Eukaryota; Liliopsida;
Magnoliophyta; Liliopsida;
NCBI_TaxID=4577;
 Q9M4X4;
 Cicer arietinum (Chickpea) (Garbanzo).
Eukaryota; Viridiplantae; Embryophyta; Tracheophyta; Spermatophyta;
Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids I;
 EMBL; AJ275305; CAB61
INTERPRO; IPR001753;
PFAM; PF00378; ECH; 1
SEQUENCE 264 AA; 2
 "Jittery, a low-copy, Mu-related BSMV infection.";
 TRANSPOSASE
 SEQUENCE FROM N.A.
STRAIN=CV. CASTELLANA; TISSUE-ETIOLATED
 PUTATIVE ENOYL COA HYDRATASE
 Q9SMK7
 Yan X., Li J., Cowperthwaite M.,
 TRANSPOSON-JITTERY;
 SEQUENCE FROM N.A.
 01-JUN-2000
 185
 Submitted (NOV-1999) to the
 An putative enoyl CoA hydratase expressed
 abrador E., Esteban R.,
 CBI_TaxID=3827;
 586 DDGNLLEEKPE 596
 1 EDGNVLKRSPE 11
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 PRELIMINARY;
 CAB61740.1;
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 Papilionoideae; Cicer
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 62.3%;
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 Dopico B.;
 Embryophyta; Tracheophyta; Poales; Poaceae; Zea.
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01-CCT-2000 (TrEMBLrel 15, Last annotation update)
HYPOTHETICAL 30.1 KDA PROTEIN.
 EMBL;
 "Structural analysis of plasmid pSN2 in Staphylococcus aureus: involvement in enterotoxin B production.";
J. Bacteriol. 149:642-649(1982).
EMBL; J01763; AAA91889.1; -
 Bacteria; Firmicutes; Bacillus/Clostridium group; Bacillus/Staphylococcus group; Staphylococcus.NCBI_TaxID-1280;
 Cerevisiae and Kluyveromyces lactis.
Genetics 148:1021-1029(1998).
EMBL; AF035007; AAC39438.1; -.
SEQUENCE 1314 AA; 146734 MW; 3C7
 Eukaryota; Fungi; As
Saccharomycetaceae;
 082741;
 082741
 SEQUENCE
 Plasmid
 SEQUENCE FROM N.A. MEDLINE-82119973; PubMed-7056699;
 Staphylococcus aureus
 01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-MAY-1999 (TrEMBLrel. 10, Last annotation update)
PLASMID PSN2 INCLUDES AN UNKNOWN GENE.
 "Theme and variation among silencing p
 Kluyveromyces lactis.
Eukaryota; Fungi; Ascomycota;
 01-MAY-2000 (TrEMBLrel. 13, Last annotation update SILENT INFORMATION REGULATOR 4.
Arabidopsis thaliana (Mouse-ear cress)
Eukaryota; Viridiplantae; Embryophyta;
 F7H19.100 OR AT4G22920
 EMBL;
 Plasmid pSN2.
 Astrom S.U., Rine J.;
 MEDLINE=98198828; PubMed=9539421;
 SEQUENCE FROM N.A.
 NCBI_TaxID=28985;
 .101
 Khan S.A., Novick R.P
 707 EDGNVSEKTPQ 717
 Local Similarity
 15
 1 EDGNVLKRSPE
 1 EDGNVLKR-----SPEL 12-
 14
 EEGNIIKRRTGALMLNPEL 119
 Similarity
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 158 AA;
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 CAA24595.1;
 11
 18182 MW;
 Kluyveromyces
 54.5%;
 61.5%;
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 Score 38;
Pred. No.
 Score 37.5;
Pred. No. 27;
 Saccharomycetes;
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 EF0426F315C6F659 CRC64;
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Tracheophyta;

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Search completed: June 4, 2001, 12:23:30 Job time: 593 sec
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Matches 7; Conservative
 Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids II;
Brassicales; Brassicaceae; Arabidopsis.
NCBI_TaxID=3702;
[1]
 SEQUENCE FROM N.A.

BU ARABIdOpsis sequencing project;
BU ARABIdOpsis sequencing project;
Submitted (MAR-2000) to the EMBL/GenBank/DDBJ databases.

EMBL; AL031018; CAA19807.1;
EMBL; AL161558; CAB79247.1;
Hypothetical protecin.

Hypothetical protecin.

BEQUENCE 268 AA; 30052 MW; 1B3F128F18EA79B2 CRC64;
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Peters S.A., van Staveren M., Dirkse W., Stiekema W., Mewes H.W.,
Lemcke K., Mayer K.F.X.;
Submitted (MAR-2000) to the EMBL/GenBank/DDBJ databases.
[3]
 SEQUENCE FROM N.A.

Peters S.A., van Staveren M., Dirkse W., Stiekema W., Bancroft I.,

Mewes H.W., Mayer K.F.X., Schueller C., Bevan M.;

Submitted (JUL-1998) to the EMBL/GenBank/DDBJ databases.
 2 DGNVLKRSPEL 12
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170 DGNLLNNYPEL 180
 60.7%;
 Score 37; DB 10;
Pred. No. 60;
1; Mismatches 3
 3
 Length 268;
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 Gaps
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GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.

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Result
No.
 Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
 Minimum DB seq length: 0
Maximum DB seq length: 2000000000
 Total number of hits satisfying chosen parameters:
 Scoring table:
 Database
 Sequence:
 Title:
 Run on:
 OM protein - protein search, using sw model
 Perfect score:
 Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
 Score
 Query
Match
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Gapop 10.0 , Gapext 0.5
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 390729 segs, 57163235 residues
 1 ELQVLGLQLPTP 12
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60
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 /SIDS6/gcgdata/geneseq/geneseqp/AA1993.DAT:
/SIDS6/gcgdata/geneseq/geneseqp/AA1994.DAT:
 'SIDS6/gcgdata/geneseq/geneseqp/AA1996.DAT:
 /gcgdata/geneseq/geneseqp/AA1980.DAT:
 DВ
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Y96183
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W00859
W80448
 SUMMARIES
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 Description
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| ļ | 526    | 377 | 377    | 347    | 347    | 100    | 1261   | 1184   | 1184   | 1098   | 363     | 329    | 195    | 195    | 73     | 8      | 264    | 178    | 124    | 108    | 100    | 87     | 649    | 174    | 164    | 498    | 498    | 422    | 380    | 338    | 374 | 374      | 344    | 377      |  |
| • |        |     |        |        |        |        | 20     |        |        | 21     | 21      |        | 21     |        | 19     | 20     | 20     | ٠.,    |        | μ.     | _      | 21     | 22     | 16     | 16     | 16     | 12     | 20     | 12     | 20     | 13  | 13       | 13     | 20       |  |
|   | 8      | 62  | 52     | R76626 | 53     | Y34819 | W30600 | R78519 | R32882 | в02009 | Y96748  | в12525 | Y84016 | W50256 | W50259 | Y06293 | Y37436 | G36067 | G25953 | G41005 | G25954 | G26029 | в65680 | R67586 | R67587 | R87089 | R13270 | W88360 | R13272 | W88361 | 55  | $\vdash$ | R22549 | . W86195 |  |
|   |        |     |        |        |        |        |        |        |        |        |         |        |        |        |        |        |        |        |        |        |        |        |        |        |        |        |        |        |        |        |     |          |        |          |  |
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## ALIGNMENTS

|   | RESULT | 360<br>JLT 1              |                                                         |
|---|--------|---------------------------|---------------------------------------------------------|
|   | Ü      | 60560                     | standard; peptide; 12 AA.                               |
|   | AC.    | W60560;                   |                                                         |
|   | XX     |                           |                                                         |
| * | TOT    | 18-AUG-1998 (f            | (first entry) .                                         |
|   | D.     | Oligopeptide fr           | from extracellular domain of CD64.                      |
|   | ×      |                           |                                                         |
|   | KW     | Extracellular domain;     | lomain; CD64; identification; antibody;                 |
|   | KW     | immunohistochemical;      | <pre>ical; immunofluorescent analysis; detection;</pre> |
|   | KW     | cell transformation;      |                                                         |
|   | XX     |                           |                                                         |
|   | SO     | Synthetic.                |                                                         |
|   | D X    | W00915933-A1              |                                                         |
|   | ×      |                           |                                                         |
|   | PD     | 16-APR-1998.              |                                                         |
|   | ×      | ,                         |                                                         |
|   | PF     | 07-OCT-1997;              | 97WO-NL00557.                                           |
|   | XX     |                           |                                                         |
|   | PR     | 08-OCT-1996;              | 96EP-0202791.                                           |
|   | PA     | (UYUT-) RIJKSUNIV UTRECHT | ITV (ITRECHT.                                           |
|   | X      | ,                         |                                                         |
|   | ΡI     | De Kruif CA, L            | Logtenberg T;                                           |
|   | ××     | ,                         |                                                         |
|   | DR     | WPI; 1998-240964/21.      | 4/21.                                                   |
|   | XX     |                           |                                                         |
|   | ΡŢ     | Identifying pep           | pecifically to protein                                  |
|   | PT     | expressing on p           | immobili                                                |
|   | Ϋ́     | oligo:peptide d           | erived from the target, useful for, e.g. identifying    |
| ÷ | Tq     | specific antibodies       | dies                                                    |

```
RESULT
R91439
 Query Match
Best Local Similarity
Trahes 12; Conserv
 밁
 Š
 01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
 forms of the same protein, including bispecific antibodies that bind to two non-overlapping epitopes on the same monomeric antigen or two epitopes on different molecules. The genes/oligonucleotides that encode selected peptides can be isolated and used for recombinant production of antibodies. Selected antibodies are used, e.g. for immunohistochemical or
 surface of a replicable display package, synthesising oligopeptides derived from the target protein on a solid phase, and testing for binding between the peptide and oligopenides. The method is used to screen large peptide libraries, especially to detect antibodies, or their fragments, that bind to cell markers or that can differentiate between different forms of the same protein, including bispecific antibodies that bind to
 Synthetic oligopeptides W60537-61 are derived from the extrace domain of CD64. They were synthesised on 25 polyethylene rods adjacent, non-overlapping sets. Peptide specific phages were i affinity binding to the CD64-covered rods. The specification of
N-PSDB; T14718
 Aruffo A,
 Cell surface antigen; cloning; immunoselection; therapy; diagnosis; vector; FcRI; Fc receptor.
 method for the identification of a peptide able to bind specifically to a target protein. The method comprises displaying the peptide on the surface of a replicable display package, synthesising oligopeptides
 WPI; 1996-200279/20
 25-FEB-1988;
 09-APR-1996.
 US5506126-A.
 Human FcRI (cDNA clone p98 product).
 R91439;
 R91439 standard; Protein;
 Sequence
 antibodies. Selected antibodies are used, e.g. immunofluorescent analysis, and also to detect
 Misc-difference
 30-OCT-1996
 Example 1; Page 29; 40pp;
 (GEHO) GEN HOSPITAL
 18-OCT-1993;
 1 ELQVLGLQLPTP 12
|||||||||||
1 elqvlglqlptp 12
 N
 Seed B;
 12
 100.0%; ilarity 100.0%; Conservative 0;
 in (anti)oncogenes
 AA,
 92US-0983647.
88US-0160416.
89US-0379076.
90US-0553759.
 93US-0139273
 88US-0160416
 Location/Qualifiers
 /note= "amino acid 58 is Leu in clone p135
translated product"
 CORP
 344
 English
 Score 60;
Pred. No.
 A
 Mismatches
 to detect cell transformation caused
 0.00013;
 DB 19;
 The specification describes
 immunotherapy;
 Length 12;
 Indels
 extracellular
 0;
 isolated
 Gaps
 by
s a
 0,
```

```
RESULT
Y96183
 B
 Qγ
 OSXWWW
 Query Match
Best Local 9
 Matches
 of the p98 product is truncated compared with those of the p135 and p90 products (see also R91438 and W00859). A fusion protein of FCRI and a receptor ligand will be helpful in increasing the potency of antibodies in therapy.
 for the Fc portion of IgG, normally located on cell surfaces of macrophages. 2 Other cDNA clones, pl35 (T14717) and p90 (T14719), coded for a variants of the FcRI sequence; the C-terminal sequence
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 panning; immunodiagnosis; diagnosis; immunotherapy; gene therapy;
immune disorder; infection; asthma; immune-complex disease;
 19-DEC-2000
 Y96183 standard; Protein; 344 AA.
 Sequence
 The amino acid sequence (R91439) of human FcRI was detd. from a cDNA clone, p98 (T14718), obtd. from a cDNA library using an immunoselection cloning method. FcRI is a high affinity receptor
 Example 10; Column 55-56; 79pp; English.
 Cloning of cDNA encoding cell surface antigen – useful for isolation of diagnostic and therapeutic proteins % \left(1\right) =\left\{ 1\right\}
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Homo sapiens
 Macrophage; FCRI; cell surface antigen; human;
 Human macrophage-specific FcRI.
 Misc-difference
 277
 Local
 1 ELQVLGLQLPTP 12
 elqvlglqlptp-288
 1 Similarity
12; Conserv
 344 AA;
 Conservative
 multiple sclerosis; parasitic disease;
 (first entry)
 /note=
 /note=
 /note=
117
 /note=
 Location/Qualifiers
/note=
126
 /note=
123
 64
 122
 /note=
 /note=
 note=
 /note=
 'note=
 'note=
 note=
 100.0%;
 encoded by
 "encoded by CAC"
 "encoded by
 "encoded by ACC"
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 "encoded
 "encoded
 "encoded by
 "encoded
 "encoded
 "encoded by
 "encoded by ACC"
 "encoded
 0;
 Score 60;
Pred. No.
 bу
 bу
 by CAA'
 ЪУ
 γď
 Mismatches
 GAA"
 ACC"
 CAG"
 CCC"
 cgg.
 TTC"
 GCA"
 TCC"
 0.0049;
 ĎВ 17;
 immunoselection;
 Length 344;
 Indels
 Fc receptor
 0;
 Gaps
```

0;

Misc-difference

0

```
01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
23-MAR-1990;
 physical selection of cells expressing the antigen by adhesion to (panning on) an antibody-coated substrate such as a culture dish. CSA nucleic acids isolated by the method of the invention, and the proteins they encode, are useful for immunodiagnostic and immunotherapeutic applications, including the diagnosis and
 The present sequence is that of a human macrophage specific FCRI, as deduced from cDNA clone pl35 (see A50631), which was isolated from a cDNA library expressed in COS cells using a novel method of
 Isolated nucleic acid molecule encoding the CD19 cell surface antigen, useful for immunodiagnosis and immunotherapy of immune-mediated infections or disorders, e.g. asthma, immune-complex disease, parasitic
 Misc-difference
 immune-complex disease, amyloidosis, parasitic diseases or multiple sclerosis. FcRI is a high affinity receptor for the Fc portion of IgG, normally located on the cell surfaces of macrophages. The
 treatment of immune-mediated infections, diseases, and disoanimals, including humans. These disorders include asthma,
 based upon transient expression of a CSA in eukaryotic cells and
 the invention designed to isolate CSA nucleic acids.
 Example
 Stamenkovic I,
 (GEHO) GEN
 28-OCT-1998;
 29-AUG-2000
 US6111093-A
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 2000-586382/55.
DB; A50631.
 10;
 interfere with such bonding, or
 HOSPITAL CORP
 Column 53-55;
 88US-0160416.
89US-0379076.
90US-0498809.
90US-0553759.
 /note=
140
 /note-
136
 92US-0983647
 /note=
332
 /note=
268
 Seed
 98US-0181612
 /note=
338
 /note=
333
 /note=
306
 /note=
305
 /note=
220
 /note=
213
 /note-
139
 /note=
134
 /note=
129
 /note=
216
 В,
 "encoded
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 English.
 CCT*
 GTG'
 CGT.
 TTC
 CAA
 AAT"
 CCA"
 GTT"
 AAT"
 GGC"
 GGT"
 AAC'
 AAT"
 diseases, and disorders in
 cause it to occur on
 The method is
```

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밁
 Q
 35×30
 Query Match
Best Local S
Matches 12
 01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
18-OCT-1993;
 for the FC portion of IgG, normally located on cell surfaces of macrophages. Another isolated cDNA clone, p90 (T14719), coded for a variant (W00859) of the FCRI sequence, and a third clone, p98 (T14718), coded for an FCRI (R91439) having a different C-terminal sequence. A fusion protein of FCRI and a receptor ligand will be helpful in increasing the potency of antibodies in therapy.
 protein of FCRI and a receptor ligand will be helpful to increase the potencies of antibodies in therapy.
 The amino acid sequence (R91438) cDNA clone, p135 (T14717), obtd.
 Example 10; Column 55-56; 79pp;
 Cloning of cDNA encoding cell surface antigen of diagnostic and therapeutic proteins
 N-PSDB;
 Aruffo A,
 09-APR-1996
 US5506126-A
 Misc-difference
 Homo sapiens
 therapy; diagnosis;
 Cell surface
 Human FcRI (cDNA clone pl35 product).
 30-0CT-1996
 R91438;
 R91438 standard;
 cDNA clone, p135 (T14717), obtd immunoselection cloning method.
 (GEHO) GEN HOSPITAL
 25-FEB-1988;
 Misc-difference
 277 elqvlglqlptp 288
 4
 1996-200279/20
DB; T14717.
 ELQVLGLQLPTP
 100.0%;
al Similarity 100.0%;
12; Conserva+*...
374 AA
 Seed B;
 344 AA;
 antigen; cloning;
 (first entry)
 92US-0983647.

88US-0160416.

89US-0379076.

90US-0553759.

93US-0139273.
 88US-0160416
 Location/Qualifiers 25
 /note=
 /note=
 Protein;
 12
 vector;
 CORP.
 "amino acid
translated
 "amino acid
translated
 374
 obtd.
 0;
 FcRI; Fc receptor.
 Score 60;
Pred. No.
 Ā.
 immunoselection;
 English
 Mismatches
 from a cDNA library using an FCRI is a high affinity receptor
 of human FcRI was detd.
 25 is Thr
 58 is Val in p90 clone product"
 0.0049;
 DΒ
 21;
 in
 0

 useful for isolation

 clone p90
 immunotherapy;
 Length
 Indels
 and
 0;
 a
 p98
 Gaps
```

Sequence

멍 δõ

W80448 standard;

Protein;

374 A

```
RESULT
W80448
 RESULT
W00859
 Query Match
Best Local s
 Matches
 Query Match
Best Local S
Matches 12
 01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
 The amino acid sequence (W00859) of human FCRI was detd. from a cDNA library using an immunoselection cloning method. FCRI is a high affinity receptor for the FC portion of IgG, normally located on cell surfaces of macrophages. Other cDNA clones (see also T14717-18) coded for variants (R91438-39) of the FCRI sequence. A fusion protein of
 277
 Sequence
 potency
 Example
 Cloning of cDNA encoding cell surface antigen - useful for isolation of diagnostic and therapeutic proteins
 (GEHO)
 18-OCT-1993;
 09-APR-1996
 US5506126-A
 Cell surface antigen; cloning;
 Human FcRI (cDNA clone p90 product).
 25-FEB-1988;
 Homo sapiens
 therapy; diagnosis;
 30-OCT-1996
 W00859;
 W00859 standard;
 1 ELQVLGLQLPTP 12
 1 ELQVLGLQLPTP 12
111111111111
277 elqvlglqlptp 288
 elqvlglqlptp 288
 J
 1996-200279/20.
DB; T14719.
 ch 100.0%;
1 Similarity 100.0%;
12; Conservative
 Ą,
 GEN
 l a receptor
of antibodié
 l Similarity
12; Conserve
 374
 antibodies
 Column 55-56; 79pp; English.
 HOSPITAL CORP
 100.0%; ilarity 100.0%; Conservative
 (first
 AA;
 88US-0160416.
89US-0379076.
90US-0553759.
93US-0139273.
 92US-0983647
 88US-0160416
 Protein;
 ligand will be helpful in
 vector; FcRI;
 entry)
 therapy.
 374
 0
 Score
Pred.
 0,
 A
 Score 60; DB 17;
Pred. No. 0.0054;
 immunoselection;
 Mismatches
 Mismatches
 FC
 No ;
 receptor.
 0.0054;
 ĎΒ
 DB 17;
 ço;
 0;
 immunotherapy;
 Length
 Indels
 Length 374;
 Indels
 the
 0;
 0;
 Gaps
 Gaps
 0;
 0;
 This is the amino acid sequence of human Fc receptor I (FCRI), as deduced from cDNA clone pl35 (see V63456) isolated using a rapid immunoselection cloning method from a cDNA library expressed in COS cells. The cDNA library was constructed from polyA RNA of cells from a single patient undergoing extracorporeal interleukin-2
 25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
21-MAY-1997;
 Example 10; Column 53-54;
 New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially human lymphocyte antigenic sequences
 WPI; 1998-609251/51.
N-PSDB; V63456.
 Aruffo A,
 (GEHO) GEN HOSPITAL CORP.
 01-DEC-1992;
 21-MAY-1997;
 03-NOV-1998.
 US5830731-A
 Misc-difference
 Misc-difference
 Misc-difference
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 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Homo sapiens
 cloning
 Fc receptor I;
 Human Fc receptor
 W80448;
 07-JUN-1999
 Seed
 92US-0983647.
88US-0160416.
89US-0379076.
 (first entry)
 90US-0553759
97US-0861205
 /note=
, 23
 В;
 97us-0861205
 /note=
256
 /note=
176
 /note=
159
 /note=
 /note=
85
 60
 /note=
 /note=
 'note-
 /note=
 Location/Qualifiers
 note=
 'note=
 note=
 /note-
 cell surface antigen; lymphocyte; human;
 "encoded by
 "encoded
 "encoded
 "encoded
 "encoded
 "encoded
 "encoded
 "encoded
 . encoded
 "encoded by
 "encoded
 "encoded by
 "encoded
 "encoded
 75pp; English.
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 ьy
 by CTG'
 by CTC"
 by TGG
 GGG"
 GTC"
 AAC"
 ATG"
 GGC"
 CAA"
 TCC"
 CCC"
 GAC"
 CCC"
 AAT"
```

ρy Дb

SQ XXX

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RESULT
W97833
 밁
 Query Match
Best Local S
Matches 12
 the isolation and cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V63442-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated genes can be expressed in a prokaryotic or eukaryotic host cells to produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V63444) which allow the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and immunotherapeutic applications, including the diagnostic and transport of immunotherapeutic applications, including the diagnostic and transport in the content of t
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Homo sapiens
 Fc receptor I;
 Human Fc receptor
 07-JUN-1999
 W97833;
 W97833 standard; Protein; 374 AA.
 Sequence
 immunotherapeutic applications, treatment of immune-mediated in
 adhesion to an antibody-coated substrate.
 expression libraries is based on transient expression of an in eukaryotic cells and selection of cells expressing the ar
 induction therapy.
 277
 Local Similarity 100 hes 12; Conservative
 1 ELOVIGIOLPTP 12
 also obtained
 374 AA;
 (first entry)
 immune-mediated infections, diseases,
 /note-
. 85
 /note=
. 77
 FCRI;
 /note-
159
 /note=
99
/note=
256
 /note=
176
 /note=
171
 103
 60
 Location/Qualifiers
 /note=
 /note=
 note-
 note
 note-
 note=
 Clones p90 (see x07372) and .A novel method for cloning
 100.0%;
 cell surface antigen; lymphocyte; human;
 "encoded
 "encoded by
 "encoded
 "encoded by
 "encoded by CTC"
 "encoded by TGG
 "encoded
 "encoded
 "encoded by
 "encoded by
 "encoded
 "encoded
 "encoded by GTC
 Score 60; DB 19;
Pred. No. 0.0054;
 уd
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 by
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 bу
 Mismatches
 . ccc.
 ATG"
 CAA.
 AAT"
 CCC.
 CTG
 GAC
 AAC
 GGC
 TCC"
 x07372) and p98 (s
for cloning cDNAs
 DB 19;
 method is useful
 Length 374;
 Indels
 and disorders of
 (see X07373)
 of an antigen by
 from mammalian
 0;
 Gaps
 for
 0;
```

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밁
 Q
 8888
 W97834
 RESULT
 Query Match
Best Local S
Matches 12
 01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
21-MAY-1997;
 genes can be expressed in a prokaryotic or eukaryotic host cells to produce the encoded protein. The invention also provides high efficiency expression vectors (see v63441 and v63444) which allow the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and
 New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially
Human Fc receptor I.
 07-JUN-1999
 W97834;
 W97834 standard; Protein; 374 AA
 were also obtained. A novel method for cloning cDNAs from mammalian expression libraries is based on translent expression of an antigen in eukaryotic cells and selection of cells expressing the antigen by adhesion to an antibody-coated substrate. The method is useful for
 This is the amino acid sequence of human Fc receptor I (FcRI), deduced from cDNA clone p90 (see x07372) isolated using a rapi
 WPI; 1998-609251/51.
N-PSDB; X07372.
 21-MAY-1997;
 03-NOV-1998
 Sequence
 animals,
 antigens from mammalian lymphocytes (see W80440-55)
 has been used to clone genes (see V63442-63) encoding cell surface
 the isolation and cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. It
 cells.
 Example 10; Column 53-54; 75pp; English.
 human lymphocyte antigenic sequences
 Aruffo A,
 (GEHO) GEN HOSPITAL CORP
 US5830731-A.
 induction therapy
 reatment of immune-mediated infections, diseases,
 from a single patient
 mmunotherapeutic applications,
 mmunoselection cloning method from a cDNA library expressed in COS
 Local Similarity hes 12; Conser
 1 ELOVIGIOLPTP 12
 The cDNA library was constructed from polyA RNA of cells
 including humans
 Seed B;
 374 AA;
 100.0%; Score 60; DB 19; ilarity 100.0%; Pred. No. 0.0054; Conservative 0; Mismatches 0
 (first entry)
 97US-0861205
 88US-0160416
89US-0379076
 90US-0553759
 9205-0983647
 97US-0861205
 /note≕ "encoded by GGG"
 ent undergoing extracorporeal interleukin-2 Clones pl35 (see V63456) and p98 (see X07373)
 including the diagnosis
 DB 19;
 isolated using a rapid
 0;
 Length 374;
 Indels
 and disorders of
 The isolated host cells to
 0
 Gaps
```

Fc receptor I; FcRI; cell surface antigen; lymphocyte; human;

<u>.</u>..

```
from a single patient undergoing extracorporeal interleukin-2 induction therapy. Clones p135 (see V63456) and p90 (see X07372) were also obtained. A novel method for cloning cDNAs from mammalian expression libraries is based on transient expression of an antigen in eukaryotic cells and selection of cells expressing the antigen by adhesion to an antibody-coated substrate. The method is useful for the isolation and cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V63442-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated
 New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially human lymphocyte antigenic sequences
 This is the amino acid sequence of human Fc receptor I (FCRI), deduced from cDNA clone p98/X2 (see X07373) isolated using a rimmunoselection cloning method from a cDNA library expressed in
 Example 10; Column 53-54; 75pp; English.
 WPI; 1998-609251/51.
N-PSDB; X07373.
 Aruffo A,
 Misc-difference
 (GEHO) GEN HOSPITAL CORP
 /note=
Misc-difference 171
 /note=
Misc-difference 159
 21-MAY-1997;
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Homo sapiens.
 The cDNA library was constructed from polyA RNA of cells
 88US-0160416.
89US-0379076.
90US-0553759.
97US-0861205.
 /note=
77
 92US-0983647
 97US-0861205
 /note=
103
 /note=
 /note=
99
 85
 45
 /note= "encoded by 23
 'note=
 'note=
 /note=
 Location/Qualifiers
 note=
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 note= "encoded by
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 GTC"
 ATG"
 CCC"
 CAA"
 TCC"
 AAT"
 CIG
 GAC"
 rapid
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밁
 Query Match
Best Local Similarity
Matches 12; Conserv
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Macrophage; FcRI; cell surface antigen; human; immunoselection; panning; immunodiagnosis; diagnosis; immunotherapy; gene therapy; immune disorder; infection; asthma; immune-complex disease; amyloidosis; multiple sclerosis; parasitic disease; Fc receptor.
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 genes can be expressed in a prokaryotic or eukaryotic host cells to produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V63444) which allow the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and immunotherapeutic applications, including the diagnosis and treatment of immune-mediated infections, diseases, and disorders of the contract of the contr
 Human macrophage-specific FcRI.
 Y96134 standard; Protein; 374 AA
 Sequence
 9
 elqvlglqlptp 288
 ELQVIGLQIPTP 12
 including humans.
 374 AA;
 Conservative
 /note=
 /note=
 /note=
116
 /note=
 /note=
 /note=
117
 /note=
 /note=
122
 /note=
82
 64
 Location/Qualifiers
 'note=
 note-
 note=
 100.0%;
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 "encoded by
 "encoded by
 "encoded
 "encoded
 "encoded
 0;
 Score 60; DB 19;
Pred. No. 0.0054;
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 Ьy
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 bу
 by ACC
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 Mismatches
 ACC"
 GCA"
 GAA"
 CAG"
 CAC"
 TTC"
 CAA"
 CCC.
 TCC"
 ACC"
 . CGG.
 19;
 0;
 Length
 Indels
 0;
 Gaps
 ಕ
```

```
physical selection of cells expressing the antigen by adhesion to
 Example 10; Column 53-55; 75pp; English
 diseases
 WPI;
 Stamenkovic I,
 01-DEC-1992;
 28-OCT-1998;
 29-AUG-2000
 US6111093-A
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 animals, including humans.
 based upon transient expression of a CSA in eukaryotic cells and
 (GEHO) GEN HOSPITAL CORP
 13-JUL-1989;
23-MAR-1990;
 Misc-difference
 Misc-difference
 13-JUL-1990
 25-FEB-1988
 2000-586382/55.
 92US-0983647.
88US-0160416.
89US-0379076.
90US-0498809.
 /note=
 /note-
, 216
 /note=
140
 90US-0553759
 98US-0181612
 /note=
332
 /note=
220
 213
 Seed
 /note=
306
 /note=
305
 /note-
 'note=
 В
 "encoded by
 "encoded by
 "encoded by AAT'
 "encoded by
 "encoded by
 "encoded by
 "encoded by
 "encoded by AAT"
 "encoded by TTC"
 These disorders include asthma,
 GGT"
 AAC
 GTG"
 CGT"
 CAA'
 CCA
```

```
(panning on) an antibody-coated substrate such as a culture dish. (SA nucleic acids isolated by the method of the invention, and the proteins they encode, are useful for immunodiagnostic and immunotherapeutic applications, including the diagnosis and treatment of immune-mediated infections, diseases, and disorders in the contract of immune-mediated infections, diseases, and disorders in the contract of immune-mediated infections, diseases, and disorders in the contract of immune-mediated infections.
 The present sequence is that of a human macrophage specific FCRI, as deduced from cDNA clone pl35 (see A50592), which was isolated from a cDNA library expressed in COS cells using a novel method of the invention designed to isolate CSA nucleic acids. The method is
 Isolated nucleic acid molecule encoding the CD19 cell surface antigen, useful for immunodiagnosis and immunotherapy of immune-mediated infections or disorders, e.g. asthma, immune-complex disease, parasitic
 Sequence
 surfaces other than macrophages, is useful protein of FCRI and a receptor ligand will
 immune-complex disease, amyloidosis, parasitic diseases or multiple sclerosis. FCRI is a high affinity receptor for the FC portion of IgG, normally located on the cell surfaces of macrophages. The ability to interfere with such bonding, or to cause it to occur on
 potencies of antibodies
 374
 8
 a receptor ligand w
tibodies in therapy.
 or to
ful in
ill be
 diseases, and disorders in
 therapy. A helpful to
 The method is
 A fusion
Query Match
Best Local Similarity
""" Conservi
 XX
```

Query Match
Best Local Similarity
Matches 12; Conser

Conservative

100.0%; Score 60; 100.0%; Pred. No. tive 0; Mismatch

Mismatches

0.0054;

Length 374; Indels

ġ,

Gaps

0;

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277

1 ELQVLGLQLPTP 12 

Conservative

0;

Mismatches

0

0;

Gaps.

Ó

100.0%;

Score 60; DB 21; Pred. No. 0.0054;

DB 21;

Length 374; Indels

RESULT, 11

1 ELQVLGLQLPTP 12

```
RESULT
Y96226
 밁
 receptor consist only of the extracellular portion of the receptor and are not glycosylated i.e. they do not have transmembrane domains or signal peptides. The recombinant proteins may be used in immunoassays to determine the immune status of patients with chronic diseases of the immune system, e.g. AIDS, systemic lupus erythematosus (SLE), multiple myeloma (MM), or rheumatoid arthritis. In addition, pharmaceutical compositions containing recombinant proteins may be used to treat or
 infection; immune response; CD64; monocyte; macrophage; neutrophil; eosinophil; HIV; IgG; immunosuppressive; antirheumatic; cytostatic; antiinflammatory; AIDS; SLE; multiple myeloma; rheumatoid arthritis;
 neutrophils and eosinophils. Upon Fc receptor activation, immune effector pathways are activated, leading to immune response. The present sequence may be modified to produce recombinant versions. The recombinant Fc
 cells expressing FC receptors. FcgammaRI molecules are expressed monocytes and macrophages, but expression can also be induced on
 FcgammaRI is also known as CD64. Fc receptors play an important role in defending the body against infections. First, pathogens are opsonised by serum immunoglobulins. The resulting complex then binds
 Novel Fc receptor lacking transmembrane domains, a signal peptide, glycosylation, useful for diagnosing and treating immune disorders
 03-DEC-1998; ·
 11-SEP-2000
 Y96226;
 Y96226 standard; Protein; 374 AA.
Sequence
 Disclosure;
 N-PSDB; A27466
 WPI; 2000-367968/32
 03-DEC-1998;
 07-JUN-2000
 EP1006183-A1
 systemic lupus erythematosus; tumour.
 Human; high affinity Fc receptor; FcgammaRI;
 Human high affinity Fc receptor, FcgammaRI.
 Homo sapiens
 rheumatoid arthritis
 (PLAC) MAX PLANCK GES FOERDERUNG WISSENSCHAFTEN
 10
374 AA;
 sequence is the human high affinity Fc receptor,
 Page 26-28;
 (first entry)
 98EP-0122969
 98EP-0122969.
 diseases,
 OF.
 60pp; English
 ¥
 allergies
 FcgammaRI molecules are expressed by
 or tumours,
 immunoglobulin;
 especially AIDS,
 FcgammaRI
 and
 ៥
 ៩
```

Вþ

302 elqvlglqlptp 313

```
Matches
 Query Match
Best Local
 vasotropic; antipsoriatic and antiangiogenic. The polynucleotides and polypeptides can be used for preventing, treating or ameliorating medical conditions and diagnosing pathological conditions. Polynucleotides, polypeptides, antibodies, agonists and antagonists from the present invention may be used to treat immune disorders by activating or inhibiting the proliferation, differentiation or mobilisation of immune cells, to treat disorders of haematopoletic cells, autoimmune disorders, allergic reactions, graft versus host disease and organ rejection, cancers, cardiovascular disorders; neurological disease and bacterial or viral infections. The peptides, nucleotides, antibodies, agonists and antagonists may be also be used in drug screens. C78449 to C78457 and B44240 represent sequences used in the exemplification of the present
 Sequence
 C77607 to C78448 encode the human cancer associated proteins given in B43398 to B44239. The proteins can have activities based on the tissues and cells the genes are expressed in Example of activities include: cytostatic; proliferative; vulnerary; immunomodulator; antidiabetic; antiasthmatic; antirheumatic; antiarthritic; antiinflammatory;
 antithyroid; antiallergic; antibacterial; antiviral; dermatological;
neuroprotective; cardiant; thrombolytic; coagulant; nootropic;
 Claim 11; Page 1739-1740; 2352pp; English.
 Novel isolated nucleic acids comprising sequences encoding useful for treating or diagnosing e.g. cancer
 Rosen CA, Ruben SM
 (HUMA-) HUMAN GENOME SCI INC
 12-MAR-1999;
 08-MAR-2000; 2000WO-US05882
 21-SEP-2000
 WO200055350-A1
 Homo sapiens
 neurological
 vasotropic; antipsoriatic; antiangiogenic; gene therapy; inrighted vasotropic; autoimmune disorder; haematopoletic cell disorder; autoimmune disorder;
 diagnosis; cytostatic; proliferative; vulnerary; immunomodulator; antidiabetic; antiasthmatic; antirheumatic; antiarthritic; antiviral; antiinflammatory; antithyroid; antiallergic; antibacterial; cardiant; dermatological; neuroprotective; thrombolytic; coagulant; nootropic;
 Human; cancer
 08-FEB-2001
 в43683;
 B43683 standard; Protein; 399 AA
 haemostatic;
 allergic reaction;
 Human cancer associated protein sequence SEQ ID NO:1128
 Local Similarity
les 12; Conserv
 immune disorder; haematopoletic cell disorder; autoimmune diso
illergic reaction; graft versus host disease; organ rejection;
naemostatic; thrombolytic; cardiovascular disorder; infection;
 399 AA;
 Conservative
 disease; drug screening.
 (first entry)
 associated gene; cancer
 100.0%;
 0;
Score 60; DB 21;
Pred. No. 0.0057;
Pred. No. 0.0057;
 antigen;
 gene therapy; inflammation;
 detection; cancer;
 Length 399;
 Indels
 peptides
0;
Gaps
```

ELQVLGLQLPTP 12 

0,

Query Match Best Local Similarity Whiches 11; Conserv

Conservative

86.7%; 91.7%;

Score 52; DB 20 Pred. No. 0.15; 0; Mismatches

DB 20;

Length 377;

0,:

Gaps

, ,

Sequence

```
PR
PR
 PFX
 PD
 EΉ
 S
 v81213) were isolated using a rapid immunoselection cloning method from a cDNA library expressed in COS cells. The cDNA library was constructed from polyA RNA of cells from a single patient undergoing extreorporeal interleukin-2 induction therapy. DNA sequence analysis revealed that the cDNAs encoded therapy. DNA sequence analysis revealed that the cDNAs encoded type I integral membrane proteins with 3 extracellular immunoglobulin domains. The invention provides a novel method for cloning cDNAs from mammalian expression libraries. This is based on the control of the con
 and molecular cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. CD40 cDNA (see V81198) is specifically claimed.
 This polypeptide human Fc receptor I (FcRI). FcRI cDNAs (see V81213) were isolated using a rapid immunoselection cloning
 transient expression of an antigen in eukaryotic cells and physica selection of cells expressing the antigen by adhesion to an antibody coated substrate. The method is useful for the isolation
 ODNA encoding human CD40 antigen - useful for cloning cDNA encoding cell surface antigens, constructing cDNA libraries, expression vectors for expression in eukaryotic cells or their fragments
 N-PSDB;
 WPI; 1999-069813/06
 Example 10; Column 55-56;
 Oquendo C,
 Allen
 13-JUL-1990,
07-JUN-1995;
 (GEHO) GEN HOSPITAL CORP.
 US5849898-A
 25-FEB-1988;
13-JUL-1989;
 01-DEC-1992;
 07-JUN-1995;
 Protein
 Fc receptor
 Human Fc receptor I.
 W86195 standard; Protein; 377
 15-DEC-1998
 Homo sapiens
 10-MAY-1999
 W86195;
 3-MAR-1990
 12
 V81213,
 Amiot M,
 Seed B,
 (first entry)
 92US-0983647.
88US-0160416.
89US-0379076.
 CD36; cell surface antigen; human; cDNA library.
 900S-0498809
900S-0553759
 95US-0485447
 95US-0485447
 /note- "this sequence contains a considerable number of differences from the sequence deduced from the DNA sequence given in
 Location/Qualifiers
 `Aruffo A, C
 deduced from the E
the specification"
 79pp; English.
 Camerini D, L
 Lauffer L;
 Į,
 FCRI CDNAs (see
 Stengelin
 the sequence
 physical
 ŝ
```

Ş

21-MAY-1992, (first entry)

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XXXX
 RESULT 14
R20811
 В
 밁
 Query Match
Best Local Similarity
Matches 10; Conserv
 This amino acid sequence was predicted from the cDNA sequence of cDNA clone p98/X2. It differs from the sequence predicted from clone p135 (see Q21178) at position 25; p135 encodes a Ser residue while clone p98/X2 predicts a Thr residue. At position 58, p90 (see Q21180) predicts Val and p98/X2 predicts Leu. Sequences predicted from all 3 clones show the typical features of a type I predicted from all 3 clones show the typical features of a type I
 R20811;
 integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane spanning domain, and a short, highly charged cytophasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and six Cys residues distributed among three C2 set ig-related domains. Cys residues distributed among three C1 set ig-related domains.
 R20811 standard;
 Sequence
 Example 10; Page 94a; 160pp; English.
 New CD53 cell surface antigen and DNA encoding immuno-therapy and diagnosis of haematopoietic
 N-PSDB;
 WPI; 1992-056864/07
 Seed B,
 13-JUL-1990;
 23-JAN-1992
 WO9201049-A.
 Homo sapiens
 Rapid immunoselection cloning technique; cell surface antigen; immunodlagnosis; high affinity receptor.
 Human macrophage-specific FcRI receptor encoded by clone p98.
 21-MAY-1992
 R22549;
 R22549 standard;
 (GEHO-) GEN HOSPITAL CORP.
 15-JUL-1990;
 277
 1 ELQVLGLQLPTP 12
{|||| ||||||||
280 elqvlqlqlptp 291
 13
 1 ELQVLGLQLP 10
 elqvlglqlp 286
 Aruffo A,
 the potency
 344
 Conservative
 90US-0553759
 90WO-US04986.
 Protein;
 Protein;
 Amiot M;
 80.0%;
 of antibodies
 344
 374 AA.
 Score 48; DB
; Pred. No. 0.7
0; Mismatches
 in therapy
 DB 13;
0.7;
 0;
 11.
 neoplasms,
 Length 344;
 Indels
 for
 0;
 Gaps
```

```
B
 \Sigma \Sigma \Sigma
 RESULT
 QY
 Query Match
Best Local Similarity
Matches 10; Conserv
 This amino acid sequence was predicted from the cDNA sequence of cDNA clone p135. It differs from the sequence predicted from clones p98/X2 and p90 (see Q21179 and Q21180, respectively) at position 25; p135 encodes a Ser residue and the other two clones predict a Thr residue. At position 58, p135 predicts Leu and p90 predicts Val. Sequences predicted from all 3 clones show the predicts Val. Sequences predicted from all 3 clones show the
 distributed among three C2 set Ig-related domains. of FcRI and a receptor ligand will be helpful to i
 typical features of a type I integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane-spanning domain, and a short, highly charged cytoplasmic domain. The extracellular portion contains six
Rapid immunoselection cloning technique; immunodiagnosis; high affinity receptor.
 Human macrophage-specific FcRI receptor encoded by clone p90.
 21-MAY-1992
 R22550 standard; Protein; 374 AA.
 Sequence
 potency of antibodies in therapy.
 New CD53 cell surface antigen and DNA encoding immuno-therapy and diagnosis of haematopoietic
 WPI; 1992-056864/07
N-PSDB; Q21178.
 23-JAN-1992
 Rapid immunoselection cloning technique; cell surface antigen; immunodiagnosis; high affinity receptor.
 potential N·linked glycosylation sites and six Cys residues
 Example 10;
 Seed B,
 13-JUL-1990;
 15-JUL-1990;
 WO9201049-A
 Homo sapiens
 Human macrophage-specific FcRI receptor encoded by clone p135.
 (GEHO-) GEN HOSPITAL CORP
 277 elqvlglqlp 286
 15
 ELQVLGLQLP 10
 Aruffo A,
 374 AA;
 Page 94a;
 Conservative
 (first entry)
 90US-0553759
 90WO-US04986
 Amiot
 80.0%;
 160pp;
 ×
 %; Score 48; DB
%; Pred. No. 0.7
0; Mismatches
 English.
 DB 13;
. 0.77;
 cell surface antigen;
 0;
 ij
 neoplasms,
 Length 374;
 Indels
 A fusion protein ncrease the
 for
 9
 Gaps
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0;

W09201049-A

```
X S S S S S S S S S S S S S X X X
 This amino acid sequence was predicted from the cDNA sequence of CDNA clone p90. It differs from the sequence predicted from Clone p135 (see Q21178) at position 25, p135 encodes a Ser residue CC and clone p90 predicts a Thr residue. At position 58, p90 predicts CC val while clones p135 and p98/X2 (see Q21179) both predict Leu. CC sequences predicted from all 3 clones show the typical features of CC atype I integral membrane protein and include a short hydrophobic CC signal sequence, a single 21-residue hydrophobic membrane-spanning CC domain, and a short, highly charged cytoplasmic domain. The CX strescellular portion contains six potential N-linked glycosylation CC sites and six Cys residues distributed among three C2 set Ig-related CC domains. A fusion protein of FCRI and a receptor ligand will be CC chapful to increase the potency of antibodies in therapy.
 Ouery Match 80.0%; Score 48; DB 13; Length 374; Best Local Similarity 100.0%; Pred. No. 0.77; Matches 10; Conservative 0; Mismatches 0; Indels
 Sequence
 Example 10; Page 94a; 160pp; English.
 WPI; 1992-056864/07.
N-PSDB; Q21180.
277 elqvlglqlp 286
 New CD53 cell surface antigen and DNA encoding it for immuno-therapy and diagnosis of haematopoietic neoplasms,
 Seed B, Aruffo A, Amiot M;
 15-JUL-1990;
 (GEHO-) GEN HOSPITAL CORP.
 13-JUL-1990;
 23-JAN-1992.
 1 ELQVIGIQIP 10
 374 AA;
 90US-0553759.
 90WO-US04986.
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Search completed: June 4, 2001, 12:13:33 Job time: 206 sec Ъ 0; Gaps

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Result
No.
 Database
 Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
 Maximum
 Sequence:
 Minimum
 Total number of hits satisfying chosen parameters:
 Searched:
 Scoring table:
 Title:
 Run on:
 OM protein
 Perfect score:
 Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed and is derived by analysis of the total score distribution.
 DB DB
 Score
 seq
 protein search, using sw
 length:
length:
 Match
 Query
 June 4, 2001, 12:14:32; Search time 58.74 Seconds (without alignments) 3.925 Million cell updates/sec
 BLOSUM62
Gapop 10.0 , Gapext 0.
 US-09-284-107-28
60
 Issued_Patents_AA: *
 185757 seqs, 19210857 residues
 GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
 ELQVLGLQLPTP 12
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 /cgn2_6/ptodata/2/iaa/backfiles1.pep:*
 Length DB
 2000000000
 SUMMARIES
 model
 474-988B-
 394-442B-2
 394-442B-9
706-741B-50
924-695A-50
706-741B-6
 74-988B-9
 185757
 Description
 Sequence
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 dd
Query Match
Best Local Similarity
Matches 8; Conserv
```

ALIGNMENTS

```
; MOLECULE TYPE: protein US-08-416-478A-6
 US-08-416-478A-6
 Sequence 6, Application US/08416478A Patent No. 5773578
 TELEFAX: (202) 737-35
INFORMATION FOR SEQ ID NO:
 GENERAL INFORMATION:
 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
 SEQUENCE CHARACTERISTICS:
LENGTH: 380 amino acid
 PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0'
FILING DATE: 08-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: YUN, Allen C.
 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
 TELEPHONE: (202) 628-5197
 TITLE OF INVENTION: TITLE OF INVENTION:
 APPLICANT: Hercend, APPLICANT: Triebel,
 CORRESPONDENCE ADDRESS
 NUMBER OF SEQUENCES:
 APPLICATION NUMBER: FILING DATE: 04-APR CLASSIFICATION: 435
 REGISTRATION NUMBER: 37,971 REFERENCE/DOCKET NUMBER: HE
 COUNTRY:
 TOPOLOGY:
 STATE:
 CITY: Washington
 STREET:
 ADDRESSEE:
 20004
 amino acids
 D.C.
 E: BROWDY AND NEIMARK, P.L.L.C. 419 Seventh Street, N.W., Suite 400
 U.S.A.
 Hercend, Thierry
 linear
 Floppy disk
 UMBER: US/08/416,478A
04-APR-1995
 New Proteins Produced By Human
Lymphocytes; DNA Sequences Encoding These Proteins And
Their Pharmaceutical And Biological Uses
 Frederic
 5:
 07/854,644
 HERCEND=1A
 #1.25
```

Conservative

65.0%; 72.7%;

Score 39; DB Pred. No. 13; 1; Mismatches

DB 1; Length 380;

2

Indels

0;

Gaps

0;

Qy

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Query Match
Best Local Similarity
Thes 8; Conserv:
 Вþ
 RESULT
 US-08-394-442B-6
 밁
 US-08-474-988B-6
 Sequence 6, Application US/08394442B Patent No. 5976877
 GENERAL INFORMATION:
 GENERAL INFORMATION:
 TITLE OF INVENTION:
 APPLICANT:
APPLICANT:
 TELEFAX: (202) 737-35
NFORMATION FOR SEQ ID NO:
 257
 MOLECULE TYPE:
 SEQUENCE CHARACTERISTICS:
 REGISTRATION NUMBER: 37,971
REFERENCE/DOCKET NUMBER: HE
TELECOMMUNICATION INFORMATION:
 FILING DATE: 08-SEP-1992
ATTORNEY/AGENT INFORMATION:
 CLASSIFICATION: 435
PRIOR APPLICATION DATA:
 SOFTWARE: Patentin Rel
CURRENT APPLICATION DATA:
 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
 PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/416,478
 OWNER OF SHAPESS:
CORRESPONDENCE: BROWDY AND NEIMARK, P.L.L.C.
ADDRESSEE: BROWDY AND NEIMARK, P.L.L.C.
ADDRESSEE: BROWDY AND NEIMARK, P.L.L.C.
ADDRESSEE: BROWDY AND NEIMARK, P.L.L.C.
 APPLICANT:
 APPLICANT:
 TITLE OF INVENTION:
 TOPOLOGY: linear
 APPLICATION NUMBER:
 TELEPHONE:
 FILING DATE:
 APPLICATION NUMBER: FILING DATE: 07-JUN
 2 LQVLGLQLPTP 12
 CLASSIFICATION:
 OPERATING SYSTEM:
 257 LTVLGLEPPTP 267
 LENGTH:
 COUNTRY:
 2 LQVLGLQLPTP 12
| ||||: |||
 I ||||: |||
LTVLGLEPPTP 267
 : 380 amino acids
amino acid
 20004
 YUN, Allen C.
 D.C.
 Application US/08474988B
 Triebel,
 Conservative
 (202) 628-5197
(202) 737-3528
R SEQ ID NO: 6:
 U.S.A.
 Triebel, Frederic
 IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
 Hercend, Thierry
 protein
 04-APR-1995
 07 - JUN - 1995
Lymphocytes, DNA Sequences Encoding These Proteins And
 New Proteins Produced By Human
 Thierry
 65.0%;
72.7%;
 Frederic
 NEW PROTEINS PRODUCED BY HUMAN LYNPHOCYTES, DNA SEQUENCES ENCODING THESE PROTEINS AND THEIR PHARMACEUTICAL AND BIOLOGICAL USES
 Release #1.0,
 US 07/854,644
 US/08/474;988B
 Score 39;
Pred. No.
 HERCEND-1B
 Mismatches
 Version #1.25
 DB 2;
 Length 380
 0;
 Gaps
 0;
```

```
RESULT
 JS-08-416-478A-7
 US-08-394-442B-6
 Sequence 7, Application US/08416478A Patent No. 5773578
 Query Match
Best Local
 GENERAL INFORMATION:
 CLASSIFICATION: 435
PRIOR APPLICATION DATA:
 SOFTWARE: Patentin Rel
 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
 CORRESPONDENCE ADDRESS:
ADDRESSEE: BROWDY AND NEIMARK, P.L.L.C.
STREET: 419 Seventh Street, N.W., Suite 400
 NFORMATION FOR SEQ ID NO:
 TITLE OF INVENTION:
TITLE OF INVENTION:
 NUMBER OF SEQUENCES:
 APPLICANT:
 SEQUENCE CHARACTERISTICS:
 APPLICANT:
 MOLECULE TYPE:
 COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0 CURRENT APPLICATION DATA:
 257 LTVLGLEPPTP 267
 NAME: BROWDY, ROGER L.
REGISTRATION NUMBER: 25,618
REFERENCE/DOCKET NUMBER: HER
TELECOMMUNICATION INFORMATION:
 CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
 CORRESPONDENCE ADDRESS:
 TITLE OF INVENTION:
 FILING DATE:
 APPLICATION NUMBER:
 CITY: Washington
 APPLICATION NUMBER:
 COUNTRY :-
 STATE:
 TOPOLOGY:
 TYPE:
 APPLICATION NUMBER: FILING DATE: 24-FE
 2 LQVLGLQLPTP: 12
 LENGTH:
 STREET: 419 Seven CITY: Washington STATE: D.C.
 TELEPHONE:
 COUNTRY:
 ADDRESSEE:
 20004
 Similarity
8; Conser
 D.C.
 20004
 amino acid
 U.S.A.
 E: BROWDY AND NEIMARK, P.L.L.C.
419 Seventh Street, N.W., Suite 400
 PatentIn Rejease #1.0,
 Triebel,
 380 amino acids
 Hercend, Thierry /
 (202)
 Conservative
 U.S.A.
 PatentIn Release #1.0, Version #1.25
 (202) 628-5197
202) 737-3528
08-SEP-1992
 04-APR-1995
 protein
 24-FEB-1995
 New Proteins Produced By Human
Lymphocytes, DNA Sequences Encoding These Proteins And
Their Pharmaceutical And Biological Uses
 65.0%;
72.7%;
 Frederic
 Their Pharmaceutical And Biological Uses
 US 07/854,644
 US/08/416,478A
 US/08/394,442B
 .
6:
 Score 39;
Pred. No.
 HERCEND-2A
 Mismatches
 Version #1.25
 DΒ
 Length 380;
 Indels
 Gaps
 0,
```

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REFERENCE/DOCKET NUMBER: HE TELECOMMUNICATION INFORMATION:

REGISTRATION NUMBER:

37,971

HERCEND-1A

0;

ATTORNEY/AGENT INFORMATION:

YUN, Allen

```
US-08-474-988B-7
 US-08-416-478A-7
 Sequence 7, App
Patent No. 5874
 Query Match
Best Local Similarity
 Matches 8; Conservative
 GENERAL INFORMATION:
 TELEFAX: (202) 737-3528 NFORMATION FOR SEQ ID NO:
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 FILING DATE: 04-APR-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 07/854,644
FILING DATE: 08-SEP-1992
ATTORNEY/AGENT INFORMATION:
 SEQUENCE CHARACTERISTICS:
 REFERENCE/DOCKET NUMBER: HE TELECOMMUNICATION INFORMATION:
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PRIOR APPLICATION DATA:
 SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
 CORRESPONDENCE ADDRESS:
ADDRESSEE: BROWDY AND NEIMARK, P.L.L.C.
 TUMBER OF SEQUENCES:
 ITLE OF
 REGISTRATION NUMBER:
 APPLICATION NUMBER: US/0 FILING DATE: 07-JUN-1995
 OPERATING SYSTEM:
 TELEFAX:
 TELEPHONE:
 APPLICATION NUMBER:
 COUNTRY:
 CITY: Washington
 STREET:
 TELEPHONE:
 2 LQVLGLQLPTP 12
 ENGTH:
 ENGTH:
 20004
 i: 470 amino acids amino acid
: 470 amino acids amino acid
 YUN, Allen C.
 D.C
 Application US/08474988B
 419 Seventh Street, N.W.,
 Triebel,
 (202)
 U.S.A.
 IBM PC compatible
 linear
 (202)
 (202) 628-5197
 2) 62
737-3528
737-7:
 628-5197
 PC-DOS/MS-DOS
 NEW PROTEINS PRODUCED BY HUMAN LYMPHOCYTES, DNA SEQUENCES ENCODING THESE PROTEINS AND THEIR PHARMACEUTICAL AND BIOLOGICAL USES
 Frederic
 Thierry
 65.0%;
72.7%;
 US 08/416,478
 US/08/474,988B
 37,971
 HERCEND=1B
 1; Mismatches
 Score 39; DB
Pred. No. 17;
 Version #1.25
 2;
 Length 470
 Indels
 0
 Gaps
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```
Query Match
Best Local Similarity
Thes 8; Conserve
 밁
 밁
 US-08-394-442B-7
 US-08-416-478A-2
 US-08-394-442B-7
 Sequence 2, Application US/08416478A Patent No. 5773578
 Matches
 Query Match
Best Local
 Sequence 7, Application US/08394442B Patent No. 5976877
 GENERAL INFORMATION:
 TELEFAX: (202) 737-35
INFORMATION FOR SEQ ID NO:
 GENERAL INFORMATION:
 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
 TITLE OF INVENTION:
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 APPLICANT:
 REFERENCE/DOCKET NUMBER: HE TELECOMMUNICATION INFORMATION:
 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,
FILING DATE: 24-FEB-19
 TITLE OF INVENTION:
TITLE OF INVENTION:
 CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
 CORRESPONDENCE ADDRESS
 229 LTVLGLEPPTP 239
 229 LTVLGLEPPTP 239
 UMBER OF SEQUENCES:
 STATE: D.C
 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
 NAME: BROWDY, Roger L. REGISTRATION NUMBER: 25,618
 TELEPHONE:
 COUNTRY:
 ADDRESSEE:
 2 LQVLGLQLPTP 12
 LENGTH:
 2 LQVLGLQLPTP 12
 Similarity
8; Conserv
 20004
 amino acid
 E: BROWDY AND NEIMARK, P.L.L.C.
419 Seventh Street, N.W., Suite 400
 470 amino acids
 Hercend,
Triebel;
 (202)
 Conservative
 U.S.A.
 Hercend,
 Conservative
 linear
 (202) 628-5197
 24-FEB-1995
 New Proteins Produced By Human
Lymphocytes, DNA Sequences Encoding These Proteins And
Their Pharmaceutical And Biological Uses
 New Proteins Produced By Human
Lymphocytes, DNA Sequences Encoding These Proteins And
Their Pharmaceutical And Biological Uses
AND NEIMARK, P.L.L.C.
 Frederic
 Thierry
 US/08/394,442B

 Mismatches

 Score 39;
Pred. No.
 HERCEND-2A
 Score 39; DB Pred. No. 17;
 Mismatches
 2
 Length 470;
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 Indels ·
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 Gaps
 Gaps
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Best Local Similarity
Best Action 8; Conserve
 US-08-474-988B-2
 equence 2, Application US/08474988B
atent No. 5874250
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MEDIUM TYPE: Floppy disk
COMPUTER: / IBM PC compatible
OPERATING SYSTEM: PC-POS/MS-DOS
SOFTMARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
 TELEFAX: (202) 737-3528
NFORMATION FOR SEQ ID NO:
 ENERAL INFORMATION:
 TITLE OF INVENTION:
TITLE OF INVENTION:
 ORRESPONDENCE ADDRESS:
ADDRESSEE: BROWDY AND NEIMARK, P.L.L.C.
STREET: 419 Seventh Street, N.W., Suite 400
 APPLICANT:
 NUMBER OF SEQUENCES:
 APPLICANT:
 SEQUENCE CHARACTERISTICS:
 CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/854,644
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 230 LTVLGLEPPTP 240
 FILING DATE: 08-SEP-1992
ATTORNEY/AGENT INFORMATION:
 OLECULE TYPE: protein
 TELECOMMUNICATION INFORMATION:
APPLICATION NUMBER: US/08/474,988B FILING DATE: 07-JUN-1995
 CITY: Washington STATE: D.C.
 LOCATION: 1
OTHER INFORMATION:
OTHER INFORMATION:
 ZIP: 20004
 COUNTRY:
 2 LQVLGLQLPTP 12
 NAME: YUN, Allen C.
REGISTRATION NUMBER: 37,971
REFERENCE/DOCKET NUMBER: HE
 FILING DATE:
 APPLICATION NUMBER:
 NAME/KEY:
 TOPOLOGY:
 TELEPHONE:
 OPERATING SYSTEM:
 ZIP: ~20004
 ENGTH:
 D.C.
 Washington
 amino acid
 D.C
 419 Seventh Street, N.W., Suite 400
 471 amino acids
 U.S.A.
 Conservative
 Triebel, Frederic
 Hercend, Thierry
 U.S.A.
 Peptide
 Linear
 E: Floppy disk
IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
 (202) 628-5197
 04-APR-1995
 NEW PROTEINS PRODUCED BY HUMAN LYMPHOCYTES, DAA SEQUENCES ENCODING THESE PROTEINS AND THEIR PHARMACEUTICAL AND BIOLOGICAL USES
 65.0%;
72.7%;
 US/08/416,478A
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 Score 39;
Pred. No.
 HERCEND=1A
 Mismatches
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 Indels
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US-08-394-442B-2
 JS-08-474-988B-2
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Best Local Similarity
TELEPHONE: (202) 628-5197
TELEFAX: (202) 737-3528
INFORMATION FOR SEQ ID NO: 2:
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 ATTORNEY/AGENT INFORMATION:
 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
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 NFORMATION FOR SEQ ID NO:
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CORRESPONDENCE ADDRESS
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 APPLICANT:
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LENGTH: 471 amino acids
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 PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/854,644

FILING DATE: 08-SEP-1992
 230 LTVLGLEPPTP 240
 ATTORNEY/AGENT INFORMATION:
 NAME: BROWDY, Roger L.
REGISTRATION NUMBER: 25,618
 FILING DATE: 24
CLASSIFICATION:
 PRIOR APPLICATION DATA:
 SOFTWARE:
 COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
 COMPUTER:
 COUNTRY:
 STREET:
 ADDRESSEE:
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OTHER INFORMATION:
 NAME/KEY:
LOCATION:
 NAME: YUN, Allen C.
REGISTRATION NUMBER: 37,971
 TOPOLOGY:
 APPLICATION NUMBER:
 CLASSIFICATION:
 INFORMATION:
 20004
 Washington
 D.C.
 amino acid
 Application US/08394442B
 DATE:
 E: BROWDY AND NEIMARK, P.L.L.C. 419 Seventh Street, N.W., Suite 400
 U.S.A.
 Hercend, Thierry
Triebel, Frederic
 Conservative
 PatentIn Release #1.0, Version #1.25
 (202)
 Peptide
 linear
 Floppy disk
 UMBER: US/08/394,442B
24-FEB-1995
 protein
 04-APR-1995
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1 737-3528
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 New Proteins Produced By Human
Lymphocytes, DNA Sequences Encoding These Proteins And
Their Pharmaceutical And Biological Uses
 65.0%;
72.7%;
 /note= "Hydrogen is present at the N-terminus"
 US 08/416,478
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 US-08-394-442B-2
 TELEX: 248425 EMBON NFORMATION FOR SEQ ID NO:
 APPLICATION NUMBER: FR 94/05643
FILING DATE: 06-MAY-1994
PRIOR APPLICATION DATA:
 CLASSIFICATION: 536 PRIOR APPLICATION DATA:
 SEQUENCE CHARACTERISTICS
 TELEFORMUNICATION INFORMATION:
TELEPHONE: 703-521-2297
TELEFAX: 703-685-0573
 FILING DATE: 05-MAY-1995
ATTORNEY/AGENT INFORMATION:
NAME: PATCH, Andrew J.
REGISTRATION NUMBER: 32,925
 SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/737,271
 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
 CORRESPONDENCE ADDRESS:
ADDRESSEE: YOUNG & THOMPSON
STREET: 745 South 23rd Street
OLECULE TYPE:
 2 LQVLGLQLPTP 12
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230 LTVLGLEPPTP 240
 *OLECULE TYPE: protein
 SEQUENCE CHARACTERISTICS:
 APPLICANT:
 UMBER OF SEQUENCES:
 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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 APPLICATION NUMBER: PCT/FR95/00593 FILING DATE: 05-MAY-1995
 FILING DATE:
 NAME/KEY:
 TOPOLOGY:
 REFERENCE/DOCKET NUMBER:
 COUNTRY:
 CITY: Arlington
 LOCATION:
 TOPOLOGY:
 ENGTH:
 TLE OF INVENTION:
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 2220
 : 471 amino acids amino acid
 amino acid
 Virginia
 Application US/08737271
 476 amino acids
 Conservative
 TRIEBEL,
 linear
 United States of America
 HERCEND, Thierry
 HUARD, Bertrand
 FAURE, Florence
 Peptide
 linear
 single
 protein
 24-DEC-1996
 LAG-3 PROTEIN; PRODUCTION ANTI-IDIOTYPE ANTIBODIES
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72.7%;
 SOLUBLE POLYPEPTIDE FRACTIONS OF THE LAG-3 PROTEIN; PRODUCTION METHOD, THERAPEUTIC COMPOSITION,
 /note= "Hydrogen is present at the
N-terminus"
 BET 96/844
 Score 39;
Pred. No.
 2
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 Length 471;
 Indels
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 Gaps
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Query Match
Best Local Similarity
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Best Local Similarity
Thehes 8; Conserve
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 US-09-058-555-1
 Sequence 1,
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 GENERAL INFORMATION:
APPLICANT: FAURE, Florence
 SEQUENCE CHARACTERISTICS:
 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/058,555
 FILING DATE: 06-MAY-1994
PRIOR APPLICATION DATA:
APPLICATION UNMER: PCT/FR95/00593
FILING DATE: 05-MAY-1995
ATTORNEY/AGENT INFORMATION:
NAME: DATE: 05-MAY-1995
 NAME: PATCH, Andrew J.
REGISTRATION NUMBER: 32,925
REFERENCE/DOCKET NUMBER: BETELECOMMUNICATION INFORMATION:
 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
 TITLE OF INVENTION:
 PRIOR APPLICATION DATA:
APPLICATION NUMBER: |
FILING DATE: 24-DEC-
 TITLE OF INVENTION:
 235 LTVLGLEPPTP 245
 TELEFAX: 703 C. TELEFAX: 703 C. TELEFAX: 703 CENBON
 COUNTRY: U. 22202
 TOPOLOGY:
 STRANDEDNESS:
 STREET: 745 Sout
CITY: Arlington
STATE: Virginia
 TELEPHONE:
 2 LQVLGLQLPTP 12
 LENGTH: 476 amino acids
 APPLICATION NUMBER:
 CLASSIFICATION:
 FILING DATE:
 DDRESSEE:
 No. 614327
 ESPONDENCE ADDRESS
 TYPE:
 Application US/09058555
 745 South 23rd Street
 Conservative
 703-685-0573
 United States of America
 TRIEBEL, Frederic
 Conservative
 HERCEND, Thierr
HUARD, Bertrand
 703-521-2297
 linear
 YOUNG & THOMPSON
 24-DEC-1996
 single
 Thierry
 65.0%;
72.7%;
 ANTI-IDIOTYPE ANTIBODIES
 SOLUBLE POLYPEPTIDE FRACTIONS OF THE
 LAG-3 PROTEIN; PRODUCTION METHOD,
 FR 94/05643
 US/08/737,271
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 Score 39; DB Pred. No. 17;
 DB 4; Length 476;
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 Indels
 Indels
 THERAPEUTIC COMPOSITION,
 0;
 0
 Gaps
 Gaps
 0
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235 LTVLGLEPPTP 245

2 LQVLGLQLPTP 12 | ||||:·|||

D.C

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Query Match
Best Local Similarity
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 US-08-474-988B-9
 Sequence 9, Application US/08474988B Patent No. 5874250
 Sequence
 SENERAL INFORMATION:
 -08-416-478A-9
 CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07,
FILING DATE: 08-SEP-1992
ATTORNEY/AGENT INFORMATION:
 NUMBER OF SEQUENCES: 1
 TITLE OF INVENTION:
TITLE OF INVENTION:
 APPLICANT:
 TELEFAX: (202) 737-3528

FORMATION FOR SEQ ID NO:
 tent No. 5773578
 257 LTVLGLEPPTP 267
 MOLECULE TYPE:
 SEQUENCE CHARACTERISTICS:
 REGISTRATION NUMBER: 37,971
REFERENCE/DOCKET NUMBER: HE
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 628-5197
 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 04-APR-19
 COMPUTER READABLE FORM:
 TITLE OF INVENTION:
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 CORRESPONDENCE ADDRESS:
 STREET:
 ADDRESSEE:
 STRANDEDNESS:
TOPOLOGY: 111
 STREET: 419 Seven CITY: Washington STATE: D.C.
 2 LQVLGLQLPTP 12
| ||||: |||
 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
 LENGTH:
 MEDIUM TYPE:
 COUNTRY:
 ADDRESSEE:
 INFORMATION:
Washington
 amino acid
 20004
 E: BROWDY AND NEIMARK, P.L.L.C.
419 Seventh Street, N.W., Suite 400
 YUN, Allen C.
 Application US/08416478
 Triebel,
 498 amino acids
 E: BROWDY AND NEIMARK, P.L.L.C.
419 Seventh Street, N.W., Suite 400
 U.S.A.
 Triebel,
 SS: single
linear
 Hercend, Thierry
 PatentIn Release #1.0, Version #1.25
 protein
 Floppy disk
 04-APR-1995
 65.0%;
72.7%;
 Frederic

NEW PROTEINS PRODUCED BY HUMAN

LYMPHOCYTES, DNA SEQUENCES ENCODING THESE PROTEINS

THEIR PHARMACEUTICAL AND BIOLOGICAL USES
 Thierry
 New Proteins Produced By Human
Lymphocytes, DNA Sequences Encoding These Proteins And
Their Pharmaceutical And Biological Uses
 Frederic
 US 07/854,644
 US/08/416;478A
 9:
 HERCEND-1A
 Score 39;
 Pred. No.
 Mismatches
 DB 1;
 Length 498;
 Indels
 Gaps
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밁
 Query Match
Best Local Similarity
Whiches 8; Conserve
 US-08-394-442B-9
 RESULT 14
 Sequence 9, Application US/08394442B Patent No. 5976877
 GENERAL INFORMATION:
 APPLICANT:
 TELEPHONE: (202) 628-5197
TELEPAX: (202) 737-3528
NFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
 SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
ATTORNEY/AGENT INFORMATION:
 CORRESPONDENCE ADDRESS:
 TITLE OF INVENTION:
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 NUMBER OF SEQUENCES:
 ZIP: AUVU-
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
 TOPOLOGY: 1: MOLECULE TYPE:
 REFERENCE/DOCKET NUMBER: HE TELECOMMUNICATION INFORMATION:
 APPLICATION NUMBER: US 0
FILING DATE: 08-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: YUN, Allen C
 PRIOR APPLICATION DATA:
US 08/416,478
 CLASSIFICATION: 435
PRIOR APPLICATION DATA:
 SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/394,442B FILING DATE: 24-FEB-1995 CLASSIFICATION: 424
 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
 257 LTVLGLEPPTP 267
 STREET:
 COUNTRY:
 ADDRESSEE:
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 REGISTRATION NUMBER: 37,971
 APPLICATION NUMBER: US/08/474,988B FILING DATE: 07-JUN-1995
 FILING DATE: 04-APR-1995
 OPERATING SYSTEM:
SOFTWARE: Patentr
 ZIP: 20004
 ENGTH:
 COUNTRY:
 20004-
 Washington : D.C.
 amino acid
 498 amino acids
 U.S.A.
 3: BROWDY AND NEIMARK, P.L.L.C.
419 Seventh Street, N.W., Suite
 Triebel,
 Hercend,
 Conservative
 U.S.A.
 linear
 protein
 single
 New Proteins Produced By Human
 Thierry
 65.0%;
 Their Pharmaceutical And Biological Uses
 Lymphocytes, DNA Sequences
 US 07/854,644
 Pred. No. 18;
 HERCEND-1B
 400
 Length 498;
 Encoding These Proteins And
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: LENGTH: 498 amino acids
: TYPE: amino acid
: STRANDEDNESS: Single
: TOPOLOGY: linear
: MOLECULE TYPE: protein
US-08-394-442B-9
 В
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 US-08-706-741B-50
 JS-08-706-741B-50
 Query Match
Best Local Similarity
Matches 7; Conserv
 Query Match
Best Local Similarity
Matches 8; Conserv
 Requence 50, Application US/08706741B
Vatent No. 5955593
 APPLICATION NUMBER: US/08/706,741B
FILLING DATE: 09-SEP-1996
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: HOLLAND, DONALD R.
REGISTRATION NUMBER: 35,197
REFERENCE/DOCKET NUMBER: 965017
TELECOMMUNICATION INFORMATION:
TELEPHONE: (314) 727-5188
 SENERAL INFORMATION:
 REFERENCE/DOCKET NUMBER: HERCEND-2A TELECOMMUNICATION INFORMATION: TELEPHONE: (202) 628-5197 TELEFAX: (202) 737-3528 [INFORMATION FOR SEQ ID NO: 9: SEQUENCE CHARACTERISTICS: LENGTH: 498 unino acids
 STRANDEDNESS:
TOPOLOGY: 11:
MOLECULE TYPE:
 TELEFAX: (314) 727-60
NFORMATION FOR SEQ ID NO:
 TITLE OF INVENTION: BH
NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
 SEQUENCE CHARACTERISTICS:
LENGTH: 73 amino acids
 COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC COMPATIBLE

OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
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| ||||: |||
257 LTVLGLEPPTP 267
1 ELQVLGLQLP 10
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34 ELEVLGRELP 43
 COUNTRY: UZIP: 63146
 ADDRESSEE: HOWELL & HAFERKAMP, L.C. STREET: 7733 FORSYTH BLVD., SUITE 1400 CITY: ST. LOUIS
 NAME: BROWDY, ROGER L. REGISTRATION NUMBER: 2
 amino acid
 MISSOURI .
 USA
 Conservative
 KORSMEYER, STANLEY J.

ENTION: BH3 INTERACTING DOMAIN DEATH AGONIST
 Conservative
 linear
 protein
 4) 727
4 727-6092
NO: 50:
 65.0%;
72.7%;
 60.0%;
 25,618
 Score 36; DB Pred. No. 7.3 2; Mismatches
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 US-09-284-107-28
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Copyright (c) 1993 - 2000 Compugen Ltd.
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Opaque-2 prott
regulatory pro
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Fc gamma
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 LAG-3 protein male-enhanced
 male-enhanced
 Description
adenylylcyclase ty
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 hypothetical
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 gamma (IgG
gamma (IgG
gamma (IgG
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|   |                    |                   |                    |                    |                    |                    |                 |                    |                    |                    |                    |                    |                    | •                  |                    |                    |
|---|--------------------|-------------------|--------------------|--------------------|--------------------|--------------------|-----------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|
|   | <u>.</u> 45        | 44                | 43                 | .42                | 41                 | 40                 | 39              | 38                 | 37                 | 36                 | 35                 | 34                 | 33                 | 32                 | 31                 | 30                 |
|   | S.                 | 35                | 35                 | 35                 | <u>ن</u><br>5      | 35                 | 35              | 35.                | 35                 | 35                 | 35                 | 35                 | 35                 | 35                 | 35                 | 36                 |
|   | 58.3               | 58.3              | 58.3               | 58.3               | 58.3               | 58.3               | 58.3            | 58.3               | 58.3               | 58.3               | 58.3               | 58.3               | 58.3               | 58.3               | 58.3               | 60.0               |
| • | 553                | 550               | 459                | 448                | 410                | 399                | .397            | 388                | 351                | 351                | 342                | 273                | 233                | 215                | 208                | 1264               |
|   | N                  | -                 | Ν                  | 2                  | N                  | _                  | N               | ν                  | N                  | 2                  | Ν                  | N                  | N                  | N                  | ν                  | Ν                  |
|   | S06345             | S14048            | F71257             | C56711             | T37078             | DXBHZ              | T06183          | D75496             | E71935             | C64646             | A83263             | C33282             | C72105             | T47380             | T13515             | S41603             |
|   | cell fusion glycop | RNA helicase dbp2 | hypothetical prote | casein kinase I (E | probable Na+/H+ an | protein Z4 - barle | serpin - barley | aspartate transami | dihydroorotate deh | dihydroorotate deh | dihydroorotate deh | DNA-binding protei | conserved hypothet | hypothetical prote | hypothetical prote | type V adenylyl cy |
|   |                    |                   | •                  |                    |                    |                    |                 |                    |                    |                    |                    |                    |                    |                    |                    |                    |

## ALIGNMENTS

Fc gamma (19G) receptor I (high affinity) form b - human N;Alternate names: CD64 C;Species: Homo sapiens (man) C;Date: 03-Apr-1992 #sequence\_revision 03-Apr-1992 #text\_change 21-Jan-2000 C;Accession: A41357; S03019

R;Allen, J.M.; Seed, B.
Science 243, 378-381, 1989
Science 243, 378-381, 1989
A;Title: Isolation and expression of functional high-affinity Fc receptor complementa A;Reference number: A41357; MUID:89100284
A;Accession: A41357
A;Molecule type: mRNA
A;Residues: 1-344 <ALL1>

RESULT A41357

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| A, MOIECUIE CYPE: DNA                                                                  | lase ty |
| A) ACCESSION: A3907                                                                    | proce   |
| A; Reference number: A398/8; MULD:91302383                                             | lase cy |
| A; Title: Gene organization of the human high affinity receptor for IgG, FcgammaRI (CD | prote   |
| J. Biol. Chem. 266, 13449-13455, 1991                                                  | protein |
| R; van de Winkel, J.G.J.; Ernst, L.K.; Anderson, C.L.; Chiu, I.M.                      | otein - |
| C; Accession: A39878; I70304; B41357; S03018; I57525                                   | otein - |
| C;Date: 30-Dec-1991 #sequence_revision 06-Sep-1996 #text_change 21-Jul-2000            | otein - |
| C; Species: Homo sapiens (man)                                                         | lated p |
| N;Alternate names: CD64                                                                | ohol d  |
| Fc gamma (IgG) receptor I-A (high affinity) precursor - human                          | cotein  |
| A39878                                                                                 | otein   |
| RESULT 2                                                                               | prote   |
|                                                                                        | 16.34   |
|                                                                                        | prote   |
| Db 277 ELQVLGLQLPTP 288                                                                | prote   |
|                                                                                        | on regu |
| CY I BUCKERQUETE IZ                                                                    | Teorids |
|                                                                                        | prote   |
| Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;                            | nthase  |
| .0%; Pred. No. 0.0015;                                                                 | prote   |
| 100.0%;                                                                                | ed anti |
|                                                                                        | ed anti |
|                                                                                        | ed anti |
| F;11/-1/U/Domain: immunogiobutin homotogy < MM>                                        | in prec |
| C; Keywords: alternative splicing; immunoglobulin receptor; transmembrane protein      | G) rec  |
| C;Superfamily: Fc gamma receptor I; immunoglobulin homology                            | G) rec  |
| .A; Note: the authors translated the codon ACT for residue 25 as Ala                   | JG) rec |
| A;Cross-references: EMBL:X14355; NID:g31333; PIDN:CAA32536.1; PID:g31334               |         |
| A; Residues: 1-344 <all2></all2>                                                       |         |
| A: Molecule type: mRNA                                                                 |         |
| A; Accession: S03019                                                                   |         |
|                                                                                        |         |
| A; Title: Nucleotide sequence of three cDNAs for the human high affinity Fc receptor ( | -       |
| Nucleic Acids Res. 16, 11824, 1988                                                     |         |
| R; Allen, J.M.; Seed, B.                                                               | ited,   |
| A;Cross-references: GB:X14355; GB:M21090; NID:q31333; PIDN:CAA32536.1; PID:q31334      | •       |
| D'INCOTAGES: T CAA SERVICE                                                             |         |

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A;Reference number: I55577;
A;Accession: I55577
 J. Clin. Invest. 90, 210
A; Title: Novel Fc gamma
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C;Date: 02-Jul-1996 #sequence_revision
C;Accession: I55577; I70303
 R;Porges, A.J.; Redecha, P.B.; Doebele, R.;
J. Clin. Invest. 90, 2102-2109, 1992
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A; Cross references: EMBL:X14356; NID:931331;
R; Perez, C.; Wietzerbin, J.; Benech, P.D.
Mol. Cell. Biol. 13, 2182-2192, 1993
 A;Title: Nucleotide sequence of three cDNAs
A;Reference number: S03018; MUID:89098339
A;Accession: S03018
 A; Cross-references: GB:M63830; R; Porges, A.J.; Redecha, P.B.;
 ;117-170/Domain: immunoglobulin homology <IMM2>
;293-313/Domain: transmembrane #status predicted <TMM>
;59,78,152,159,163,195,240/Binding site: carbohydrate
 Nucleic Acids Res.
 Science 243, 378-381,
 ; Map position:
 Matches
 Query Match
Best Local :
 Superfamily: Fc gamma receptor I; immunoglobulin homology Keywords: alternative splicing; glycoprotein; immunoglobu 1.15/Domain: signal sequence #status predicted <SIG>
 Cross-references:
 Gene: GDB:FCGR1A; CD64
 Molecule type: DNA
 Status: translated from GB/EMBL/DDBJ
 Reference number: 157525; MUID:93204964 Accession: 157525
 16-292/Domain: extracellular #status predicted <EXT>
 Cross-references:
 Residues:
 Molecule type: mRNA
Residues: 1-24,'S',
 Allen, J.M.;
 Title: Isolation and expression of functional high-affinity Reference number: A41357; MUID:89100284 Accession: B41357
 Title: Novel Fc gamma
 Title:
 Cross-references: GB:L03418; NID:g184840; Allen, J.M.; Seed, B.
 Accession:
 Residues:
 Molecule type: mRNA
 olecule type: mRNA
esidues: 1-24,'S',26-337,'T',339-374 <ALL>
ross-references: GB:X14356; GB:M21091; NID:g31331; PIDN:GAA32537.1;
 Reference number:
 277 ELQVLGLQLPTP 288
 1 ELQVLGLQLPTP 12
 preliminary; translated from GB/EMBL/DDBJ
 translated from GB/EMBL/DDBJ
 l Similarity
12; Conserv
 1-8 <RE2>
 cis-DNA elements
 1-374 <RES>
 receptor I-B splice form 1 precursor - nes: CD64 ,
 Seed, B.
Res. 16, 11824, 1988
 1q21-1q21
 90, 2102-2109,
 GDB:135911; OMIM:146760
 GB:S57204;
 1989
 100.0%;
 receptor I family 77; MUID:93055454
 eptor I family MUID:93055454
 involved in myeloid-cell-specific
 NID: g298692; PIDN: AAD13887.1; PID: g4261587
 Doebele,
 GB:M63835; NID:g180277; Doebele, R.; Pan, L.C.;
 0;
 Score 60;
Pred. No.
 Mismatches
 02-Jul-1996 #text_change 21-Jan-2000
 splice form
 gene
 Pan,
 gene products in
 DB 1;
0.0017;
 for
 PIDN:AAA36049.1;
 PIDN:CAA32537.1;
 products
 L.C.;
 immunoglobulin
 the human high affinity Fc receptor
 0;
 Length 374;
 Salmon, J.E.; Kimberly, R.P.
 Indels
 (Asn)
 ä
 PIDN:AAA35678.1;
Salmon, J.E.; Ki
 human
 (covalent)
 PID:g292169
 PID: 931332
 FC
 receptor;
 mononuclear cells
 0;
 expression
 receptor complementary
 mononuclear cells
 Gaps
 .1; PID:g180279
Kimberly, R.P.
 #status
 PID: 931332
 transmembrane
 and gamma
 0;
 predicto
 (FCI
 in
 Proc.
A; Reference number:
 R; Lau,
 C; Accession: B34421
 밁
 B34421
 RESULT
 C; Keywords:
 A; Gene:
 C; Genetics:
 A; Residues:
 A; Molecule type: mRNA
 B
 A; Accession: S11246
 R; Triebel,
 Qγ
 A; Map position:
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 A;Accession: I70303
 Accession: S11246
 Matches
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 Matches
 Query Match
 Exp. Med.
 257
 186
 Local
 171,
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C;Species: Mus musculus (house mouse)
C;Date: 15-Jun-1990 #sequence_revision
 A; Title: Male-enhanced antigen
 male-enhanced antigen - mouse
 A; Map position: 12p13.3-12p13.3
C; Superfamily: human LAG-3 prot
 A; Cross-references: GDB:127449;
 A;Cross-references: EMBL:x51985
A;Note: the author translated the
 A;Reference number: S11246; MUID:90237736
 A; Title: LAG-3, a novel l
 C;Species: Homo sapiens (man)
C;Date: 18-Feb-1994 #sequence_revision 10-Nov-1995 #text_change 04-Mar-2000
 C;Superfamily: Fc gamma receptor I; immunoglobulin homology C;Keywords: alternative splicing; glycoprotein; immunoglobu F;117-170/Domain: immunoglobulin homology <IMM>
 A;Residues: 1-280 <RES>.
A;Cross-references: GB:L03419; NID:g182460; PIDN:AAA35825.1;
A;Note: splice form B1
 A; Experimental source: mononuclear A; Note: splice form B2
 A; Molecule type: mRNA
A; Residues: 1-10,103-153,'A',155-280
 A; Status: preliminary; translated from
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 Cross-references: GB:L03420;
 Natl. Acad. Sci. U.S.A. 86,
 protein precursor
 2 LQVLGLQLPTP
 4 VLGLQLPTP
 preliminary
 VKGLQLPTP 194
 transmembrane protein
 1-498 <TRI>
 Similarity
8; Conserv
 Similarity 88.88; Conservative
 Jitsukawa, S.; Baixeras,
71, 1393-1405, 1990
 Conservative
 1p12-1p12
antigen gene is phylogenetically A34421; MUID:90046817
 GDB:135923;
 12
 human
 66.7%;
 protein
 86, 8462-8466,
gene is phylon
 not bind monomeric
 NID: g182461;
 Score 39; DB
Pred. No. 19;
1; Mismatches
 OMIM: 601502
 codon
 Score 40; DB
Pred. No. 6.5;
0; Mismatches
 0;
 09-Oct-1992
 CCA
 压.
 GB/EMBL/DDBJ
 for
 PIDN: AAA35826.1;
 DB
 2
 residue
 2;
 immunoglobulin receptor;
 #text_change
 IgG with high affinity.
 Length 498
 conserved
 Genevee, C.; Viegas-Pequ
 PID:g292024
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 05-Nov-1999
 and
 0;
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hypothetical protein XF2283 [imported] - Xylella fastidiosa (strain 9a5c) C:Species: Xylella fastidiosa C:Date: 18-Aug-2000 #sequence_revision 20-Aug-2000 #text_change 20-Aug-20
 В
 Q
 C;Species: Homo sapiens (man)
C;Date: 15-Jun-1990 #sequence_revision 15-Jun-1990 #text_change
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 Q
 A;Accession. A;Accession. A;Accession. A;Accession. A;Accession. A;Accession. A;Cross-references: EMBL:D17340; NID:g560052; PIDN:BAA04158.1;A;Cross-references: EMBL:D17340; NID:g560052; PIDN:BAA04158; NID:g560052; NID:g560052; NID:g560052; NID:g560052; NID:g560052; NID:g560052; NID:g560052; NID:g560052; NID:g
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 B82577
 RESULT
 Biochim. Biophys. Acta 1216, 483-486, 1993
A;Title: Cloning and sequence analysis of cDNA encoding the
A;Reference number: S39790; MUID:94092744
 male-enhanced antigen - bovine
 A; Cross-references: GB:M27938; A; Note: the authors translated
 A; Molecule type: mRNA
A; Residues: 1-174 <LAU>
 R; Lau,
 male-enhanced antigen - human
 A;Status:
 4;Cross-references: GB:M27937; NID:g187507; PIDN:AAA36208.1; PID:g307182
 R; Kondo, M.; Sato, S.; Sutou,
 ;Lau, Y.F.C.; Chan, K.; Sparkes, R.
roc. Natl. Acad. Scl. U.S.A. 86, 8462-8466, 1989
;Title: Male-enhanced antigen gene is phylogenetically conserved
;Reference number: A34421; MUID:90046817
 Accession: A34421
 Status: preliminary;
 Species: Bos primigenius taurus (cattle);Species: 19-May-1994 #sequence_revision 19-Apr-1996 #text_change 20-Jun-2000
 Accession: A34421
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 Query Match
Best Local
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 Query Match
Best Local :
 Matches
 Query Match
Best Local
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:| || || |
100 IQALGLHLPDP 110
 89
 9
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 2 LQVLGLQLPTP 12
 2 LOVIGIOLPTP 12
 IQALGLHLPDP 99
 preliminary
 Similarity 7; Conserv
 Similarity 63.0
7; Conservative
 #sequence_revision 20-Aug-2000 #text_change 20-Aug-2000
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 63.3%;
 63.3%;
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 Score 38;
Pred. No.
 Score 38; DB 2; Pred. No. 9.6;
 Score 38; D
Pred. No. 9;
 Mismatches
 Mismatches
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 Length 185
 Length 174;
 Length 174
 Indels
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 bovine testis-derived male-
 PID:9560053
 05-Nov-1999
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 Gaps
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В
 A;Title: The genome sequence of the plant pathogen Xylella fastidiosa. A;Reference number: A82515; MUID:20365717
A;Note: for a complete list of authors see reference number A59328 below
 C;Superfamily: threonine synthase
C;Keywords: carbon-oxygen lyase; phosphoprotein;
F;105/Binding site: pyridoxal phosphate (Lys) (compared to the compared A;Authors: Gnehm, C.L.; McDonald, L.A.; Small, K.V.; Fraser, C.M.; Smith, H.O.; Vente A;Title: Whole-genome random sequencing and assembly of Haemophilus influenzae Rd. A;Reference number: A64000; MUID:95350630
A;Accession: H64047
 δõ
 J.D.; Junqueira, M.L.; Kemper, E.L.; Kitajima, J.P.; Krieger, J.E.; Kuramae, E.E.; L. chado, M.A.; Madeira, A.M.B.N.; Madeira, H.M.F.; Marino, C.L.; Marques, M.V.; Martino, A.M.B.N.; Markino, A.Y.; Markino, C.F.M.; Miracca, E.C.; Miyakino, F.G.; Nunes, L.R.; Oliveira, M.A.; de Oliveira, M.C.; de Oliveira, R.C.; Palmieri, F.G.; Nunes, L.R.; Oliveira, M.A.; de Oliveira, M.C.; de Oliveira, R.C.; Palmieri,
 A;Molecule type: DNA
A;Residues: 1-425 <TIGR>
 Science 269, 496-512, 1995
 R;Fleischmann, R.D.; Adams, M.D.; White, O.; Clayton, R.A.; Kirkness, E.F.; Ker; Gocayne, J.D.; Scott, J.; Shirley, R.; Liu, L.I.; Glodek, A.; Kelley, J.M.; V.D.M.; Brandon, R.C.; Fine, L.D.; Fritchman, J.L.; Fuhrmann, J.L.; Geoghagen,
 C; Date:
 threonine synthase (EC 4.2.99.2) - Haemophilus influenzae (strain Rd C;Species: Haemophilus influenzae
 A; Gene:
 A; Contents: annotation
 A; Reference number: A59328
 , F.G.; Nunes, L.R.; Oliveira, M.A.; de Oliveira, M.C.; de Oliveira, R.C.; Palmis
Rodrigues, V.; Rosa, A.J. de M.; de Roa Jr., V.E.; de Sa, R.G.; Santelli, R.V.;
A;Authors: da Silva, A.C.R.; da Silva, F.R.; da Silva, A.M.; Silva Jr., W.A.; da
M.; Tsuhako, M.H.; Vallada, H.; Van Sluys, M.A.; Verjovski-Almeida, S.; Vettore,
 submitted to GenBank, June 2000
A; Authors: Ferreira, V.C.A.; Ferro, J.A.; Fraga, J.S.; Franca, S.C.; Franco,
 as-Neto,
 R;Simpson, A.J.G.; Reinach, F.C.; Arruda
Briones, M.R.S.; Bueno, M.R.P.; Camargo,
 A; Experimental source:
 A; Cross-references: GB: AE004040;
 A;Residues:
 A; Status: preliminary
 A; Accession: B82577
 Nature 406, 151-157
 C; Accession:
 A;Cross_references: GB:U32694; GB:L42023; NID:g1573035; PIDN:AAC21765.1; PID:g1573038
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 Matches
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 .1 ELOVIGLOIPTP 12
3 QVLGLQLPTP 12
 18-Aug-1995 #sequence_revision 18-Aug-1995 #text_change 16-Jul-1999
 1-317 <SIM>
 Similarity 6; Conser
 H64047
 The Xylella fastidiosa Consortium
 Docena, C.
 Conservative
 193
 strain
 ; El-Dorry, H.; Facincani, A.P.; Ferreira, A.J
 63.3%;
 63.3%;
 9a5c
 Score 38; DB:
Pred. No. 24;
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 GB:AE003849; NID:g9107437;
 Arruda, P.; Abreu, F.A.; Acencio, M.; Alvarenga, R. amargo, A.A.; Camargo, L.E.A.; Carraro, D.M.; Carrer
 Score
Pred.
 Mismatches
 ₩
38;
 oprotein; pyridoxal phosphate; threonine biosy (Lys) (covalent) #status predicted
 .DB
17;
 of the Organization for Nucleotide
 Ņ
 1;
 Length 425;
 Length 317;
 Indels
 L.E.A.; Carraro, Ferreira, A.J.S.
 PIDN:AAF85082.1; GSPDB:GN
 0,
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 Gaps
 KW20)
 ranco, M.C.; Fr
ramae, E.E.; La
, M.V.; Martins
.C.; Miyaki, C.
 Kerlavage
1.; Weldman
 0
 N.S.M.
 Sawa
Silv
 Seq
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386 RILGIQLPLP 395

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M: Shen, M.; Vamathevan, J.J.; Lam, S.; Smith, H.O.; Venter, J.C.; Fraser, Science 286, 1571-1577, 1999
 A; Title: Genome sequence of A; Reference number: A75250;
 transcription regulator, IclR family - C.Species: Deinococcus radiodurans C.Date: 03-Dec-1999 #sequence_revision
 R;White, O.; Eisen, J.A.; Heidelberg,
 C; Accession: C75611
 \;Status: preliminary
 밁
 Ş
 C; Superfamily: phosphoglycerate mutase; phosphoglycerate mutase homology
 Accession: E71478
 망
 Query Match
Best Local Similarity
Thes 7; Conserve
 A; Map position:
 A; Gene: CESP:C54G6.4
 R;Tin-Wollam, A.; Graves, T.; Ozersky, submitted to the EMBL Data Library, Jan
 probable phosphoglycerate mutase -
 ;Cross-references: GB:AE001343; GB:AE001273; NID:g3329176; PIDN:AAC68317.1; PID:g33291;
Experimental source: serotype D, strain UW-3/Cx
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 Genetics:
 Date: 13-Sep-1998 #sequence_revision 13-Sep-1998 #text_change 16-Jul-1999 Accession: E71478
 Residues: 1-226 <ARN>
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Experimental source: strain Bristol N2; clone:C54G6
 Molecule type: DNA
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 Status: preliminary; translated from GB/EMBL/DDB
 Reference number: Accession: T32913
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 195
 Local Similarity
nes 8; Conserv
 1 ELQVLGLQLPT 11
 98 QQLGVQLPVP 107
 mbd
 3 QVLGLQLPTP 12
 EEQVLSLELPT 205
 Caenorhabditis elegans
 Conservative
 Conservative
 61.7%;
72.7%;
 61.7%;
 MUID:20036896
 the radioresistant bacterium
 Caenorhabditis elegans (fragment)
 Score 37; DB.
Pred. No. 18;
1; Mismatches
 ۲,
 Score 37; DB Pred. No. 10; 1; Mismatches
 Chlamydia trachomatis (serotype D,
 January 1998
 J.F.;
 o
ġ
 03-Dec-1999
 Deinococcus
 McDonald,
 cosmid
 2;
 C54G6
 #text_change 17-Mar-2000
 radiodurans (strain R1)
 E.K.;
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 2;
 Length 226
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 Marathe, R.;
 Indels
 Deinococcus
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 0,
 0;
 Aravind, L.;
 humans:
 radiodurans
 Gaps
 strain
 Chlamydia tra
 UW3/Cx)
 Mitchell
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 R.J.;
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Query Match
Best Local Similarity
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A; Map position:
 submitted to the RMBL Data Library, September 1996
A;Reference number: Z19925
A;Accession: T24692
 C;Species: Caenorhabditis elegans
C;Date: 15-Oct-1999 #sequence_rev
C;Accession: T24692
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 A; Molecule type: DNA
A; Residues: 1-3212 <WIL>
 A; Status: preliminary; translated from GB/EMBL/DDBJ
 R;Dobson,
 hypothetical protein T08G11.1 -
 A; Introns: 154/1; 326/3
 R:Jones, K.; Graves, T.; Ozersky, P. submitted to the EMBL Data Library, A;Description: The sequence of C. el A;Reference number: z21420
 A; Map position: 1
 C; Genetics:
 A;Cross-references: EMBL:AF106584;
A;Experimental source: strain Brist
 A; Molecule type: DNA
A; Residues: 1-424 < JON>
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 A; Gene: CESP:F54A5.1
 A; Status: preliminary; translated from GB/EMBL/DDB.
 Query Match
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Thehas 7; Conserve
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 hypothetical protein F54A5.1 - Caenorhabditis elegans
 A;Map position: 2
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 A; Gene: DRA0152
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 Genetics:
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 Accession:
 Species:
 388 QQLGVQLPVP 397
 205 LAALGVSLPTP 215
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 3 QVLGLQLPTP 12
 2 LQVLGLQLPTP
 Caenorhabditis elegans
 149/3;
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 #sequence_revision 15-Oct-1999 #text_change 15-Oct-1999
 182/3; 527/2; 766/1; 1070/3; 1769/3; 1834/3; 2070/2;
 61.7%;
 Bristol
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 Score 37;
Pred. No.
 P.; Wilson, R.
Y, November 1998
elegans cosmid F54A5
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tol N2; clone F54;
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 Mismatches
 2;
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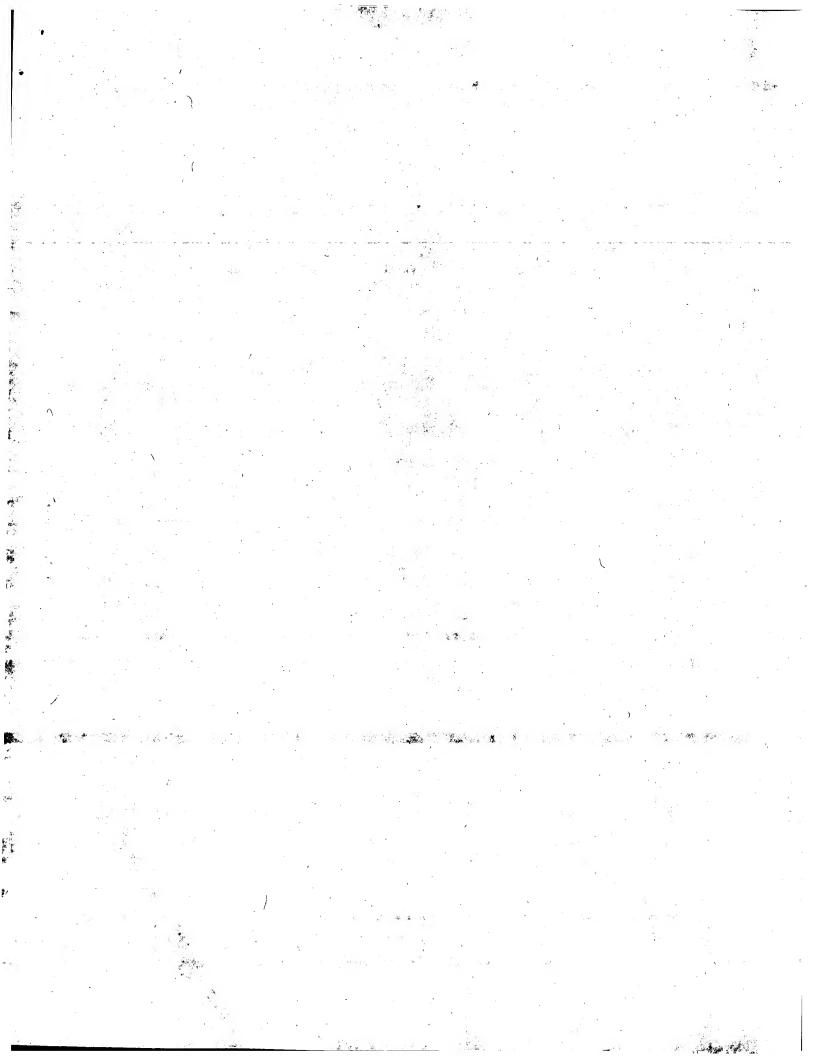
61.7%;

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A;Cross-references: EMBL:X84724; NID:g780165; PIDN:CAA59211.1; PID:g780172
A;Cross-references: EMBL:X84724; NID:g780165; PIDN:CAA59211.1; PID:g780172
R;Hess, M.; Cuzange, A.; Rulgrok, R.W.H.; Chroboczek, J;; Jacrot, B.
J. Mol. Biol. 252, 379-385, 1995
A;Title: The avian adenovirus penton: two fibres and one base.
A;Reference number: S59067; MUID:96025073
A;Contents: annotation
C;Superfamily: fowl adenovirus 1 hypothetical 16.34K protein
 밁
 R;Hess, M.; Cuzange, A.; Chroboczeck, J.; Ruigrok, R.; Jacrot, B. submitted to the EMBL Data Library, February 1995
A;Description: The sequence of the two fibers of an avian adenovirus (CELO) and organise A;Reference number: S54125
A;Accession: S54130
 ypothetical 16.34K protein - fowl adenovirus 1
;Species: Aviademovirus gall (fowl adenovirus 1, CELO)
;Date: 08-7ul-1995 #sequence_revision 14-Feb-1997 #text_change 22-Oct-1999
;Accession: S54130
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 6 GLQLPTP 12
|||:|||
39 GLQMPTP 45
 7; Conservative
 Score 36; DB 2; Length 144; Pred. No. 17; 1; Mismatches 0; Indels
 1; Mismatches _ 3; Indels
 ò
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 Gaps
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 0
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GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.

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Title:
Perfect score:
Sequence:
 Result
No.
 Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
 Minimum DB seq length: 0
Maximum DB seq length: 2000000000
 Database :
 Total number of hits satisfying chosen parameters:
 Searched:
 Scoring table:
 Run on:
 OM protein - protein search, using sw
Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
 Score
 Query
Match
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 US-09-284-107-28
60
 93435 seqs, 34255486 residues
 BLOSUM62
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 Gapop 10.0 , Gapext 0.5
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VGLF_NDVI
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RL1X_RAT
PYRD_AGRAE
FCG1_MOUSE
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ROK_RABIT
 CYA5_RABIT
CYA5_CANFA
CYA5_CANFA
PYRD_HELPJ
PYRD_HELPY
PRTZ_HORVU
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LAG3_HUMAN
MEA1_PIG
 ROK_MOUSE
KG3A_HUMAN
PT1_STAAU
 MEA1_HUMAN
MEA1_BOVIN
MEA1_MOUSE
THRC_HAEIN
P2X2_CAVPO
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BID_MOUSE
 OP2_MAIZE
NH25_CAEEL
 Ħ
 CYA5_RAT
 KC13_HUMAN
 161_HUMAN
 SUMMARIES
 mode1
 93435
 P12314 homo sapien P18627 homo sapien
 Description
 oryctolagus
mus musculu
 agrocybe ae
 newcastle d
 rattus norv
schizosacch
 caenorhabdi
 mus musculu
 homo sapien
 mus musculu
 homo sapien
 homo sapien
 helicobacte
hordeum vul
 canis famil
 oryctolagus
 rattus norv
 zea mays
 mus musculu
 saccharomyc
 cavia porce
 haemophilus
 homo sapien
 sus scrofa
 rattus
 helicobacte
 homo sapien
 norv
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|   | 45         | . 44       | 43          | 42          | 41          | 40          | 39          | 38          | 37          | 36          | 35          | 34          |   |
|---|------------|------------|-------------|-------------|-------------|-------------|-------------|-------------|-------------|-------------|-------------|-------------|---|
| • |            |            |             |             |             |             | _           |             | ٠.          |             | •           |             |   |
| • | 33         | ω<br>ω     | 33          | ω           | ω<br>ω      | 33          | ω<br>ω      | ω<br>Ψ      | ω<br>ω      | 34          | 34          | 34          |   |
| • | 55.0       | 55.0       | 55.0        | 55.0        | 55.0        | 55.0        | 55.0        | 55.0        | 55.0        | 56.7        | 56.7        | 56.7        |   |
|   | 322        | 296        | 296         | 249         | 231         | 213         | 203         | 203         | 109         | 2787        | 681         | 585         |   |
|   | 1          | ب          | -           | ۲           | -           | -           | 1           | _           | -           | ب           | بز          | 1           |   |
|   | PNUC_SALTY | SAPC_SALTY | SAPC_ECOLI  | Y768_METJA  | BID2_ECOLI  | YOUB_CAEEL  | VATD_CHLTR  | VATD_CHLMU  | YC44_METMP  | TEL1_YEAST  | BRC2_HALRO  | YAOH_SCHPO  |   |
| • |            |            |             | -           |             |             | :           |             |             |             |             |             |   |
|   | P24520     | P36669     | Q47624      | Q58178      | P77201      | P34658      | 084308      | 09pk87      | 050249      | P38110      | 001409      | 210096      |   |
|   | salmonella | salmonella | escherichia | methanococc | escherichia | caenorhabdi | chlamydia t | chlamydia m | methanococc | saccharomyc | halocynthia | schizosacch | , |

## ALIGNMENTS

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| MBL; X14356; CAA32537. MBL; X14355; CAA32536. IR; S03018; S03018. IR; S03019; S03019. IR; A41357; A41357. IR; B41357; B41357. SSP, P12319; IALT. IM; 146760; fam; PF00047; 19; 3.                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             | This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch). | -!- SUBCELLULAR LOCATION: TYPE I MEMBRANE PROTEIN!- ALTERNATIVE PRODUCTS: 2 ISOFORMS; A (SHOWN HERE) AND B; ARE PRODUCED BY ALTERNATIVE SPLICING!- TISSUE SPECIFICITY: MONOCYTE, MACROPHAGE-SPECICIC!- SIMILARITY: CONTAINS 3 IMMUNOGLOBULIN-LIKE C2-TYPE DOMAINS!- DATABASE: NAME-PROW; NOTE-CD guide CD64 entry; | SEQUENCE FROM N.A.  SEQUENCE FROM N.A.  MEDLINE-89100284; PubMed-2911749;  Allen J.M., Seed B.;  "Isolation and expression of functional high-affinity Fc receptor complementary DNAs.";  Science 243:378-381(1989).  -I- FUNCTION: BINDS TO THE FC REGION OF IMMUNOGLOBULINS GAMMA. HIGH AFFINITY RECEPTOR. | D=9606; FROM N.A. 9098339; PubMed=2974947; ., Seed B.; de sequence of three cDNAs for t (FCRI)."; cids Res. 16:11824-11824(1988). | LLT 1  LHUMAN STANDARD; PRT; 374 AA.  FCG1_HUMAN STANDARD; PRT; 374 AA.  P12314; P12315; 01-CCT-1989 (Rel. 12, Created) 01-CCT-1989 (Rel. 12, Last sequence update) 01-CCT-2000 (Rel. 40, Last annotation update) HIGH AFFINITY IMMUNOGLOBULIN GAMMA FC RECEPTOR I PRECURSOR (FC-GAMMA RI) (FCRI) (IGG FC RECEPTOR I) (CD64).  RI) (FCRI) (IGG FC RECEPTOR I) (CD64).  FCGR1A OR FCGR1 OR FCG1 OR IGFR1.  Homo saplens (Human).  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. |

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"LAG-3, a novel lymphocyte activa
J. Exp. Med. 171:1393-1405(1990).
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 MEDLINE=92364535;
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 MEDLINE=90237736; PubMed=1692078; Triebel F., Jitsukawa S., Baixera
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 Sus scrofa (Pig).
Eukaryota; Metazoa;
Mammalia; Eutheria;
 SEQUENCE FROM N.A.
 NCBI_TaxID=9823;
 MALE-ENHANCED
 EMBL; X51985; CAA36243.1;
EMBL; A21353; CAA01547.1;
PIR; S11246; S11246.
MIM; 153337; -...
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 "Male-enhanced antigen gene is phylogenetically conserved and expressed at late stages of spermatogenesis.";
Proc. Natl. Acad. Sci. U.S.A. 86:8462-8466(1989).
-I- FUNCTION: MAY PLAY AN IMPORTANT ROLE IN SPERMATOGENESIS AND/OR
 MEA1_HUMAN
Q16626;
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 Spermatogenesis; Developmental protein. SEQUENCE 172 AA; 18544 MW; 9342AFAD2C96FF25 CRC64;
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TISSUE SPECIFICITY: HIGHLY EXPRESSED IN TESTIS.
CAUTION: WAS ORIGINALLY THOUGHT TO BE THE H-Y ANTIGEN.
CAUTION: IT IS UNCERTAIN WHETHER MET-1 OR MET-11 IS THE INITIATOR.
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 "Cloning and sequence analysis of cDNA encoding derived male-enhanced antigen (Mea)."; Biochim. Biophys. Acta 1216:483-486(1993).
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01-NOV-1997 (Rel. 35, La
MALE ENHANCED ANTIGEN-1
 MEA1_BOVIN Q29407;
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DNA Seq. 6:75-85(1996).
-!- FUNCTION: MAY PLAY AN IMPORTANT ROLE IN
 MEDLINE=97063437; PubMed-8907304;
Kondo M., Terouchi S., Tsukasa N., Sato.S., Ishida N., Sutou S.;
"Genomic sequence analysis of the bovine male-enhanced antigen-1
(Mea-1) and differential localization of its transcripts and pro-
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MEDLINE=94092744; PubMed=8268232;
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 Kondo M., Sato S.,
 NCBI_TaxID=9913;
 Bos taurus (Bovine).
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 CAUTION: IT IS UNCERTAIN WHETHER MET-1 OR MET-11 IS THE INITIATOR
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MEDLINE=90046817; PubMed=2813404;
Lau Y.-F.C., Chan K., Sparkes R.S.;
"Male-enhanced antigen gene is phyLogenetically conserved expressed at late stages of spermatogenesis.";
expressed at late stages of spermatogenesis.";
expressed at late stages of spermatogenesis.";
 Bacteria; Proteobacteria; Haemophilus.
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 Spermatogenesis; Developmental protein.
SEQUENCE 174 AA; 18584 MW; C3B16361635B176F CRC64;
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 "Cloning and sequence analysis of cDNA encoding the derived male-enhanced antigen (Mea).";
Biochim. Biophys. Acta 1216:483-486(1993).

-i- EUNCTION: MAY PLAY AN IMPORTANT ROLE IN SPERMATO TESTIS DEVELOPMENT.

-i- TISSUE SPECIFICITY: HIGHLY EXPRESSED IN TESTIS.
 the European Bioinformatics Institute.
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Mammalia; Eutheria;
 Kondo M.,
 MEDLINE=94092744; PubMed=8268232;
 Mus musculus (Mouse)
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MEA1 OR MEA.
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 FOUND IN PRIMARY AND SECONDARY SPERMATOCYTES, AND SPERMATIDS, BUT THE PROTEIN ITSELF IS ONLY DETECTED IN SPERMATIS. NO EXPRESSION IN LEYDIG CELLS, SPERMATOGONIA, OR SPERM. VERY WEAK EXPRESSION IN THE HEART, KIDNEY, SPLEEN, THYMUS, AND OVARY.

DEVELOPMENTAL STAGE: EXPRESSED AS EARLY AS DAY 6 POST-PARTUM, WITH HIGHER EXPRESSION IN THE ADULT TESTIS.

CAUTION: WAS ORIGINALLY THOUGHT TO BE THE H-Y ANTIGEN.

CAUTION: IT IS UNCERTAIN WHETHER MET-1 OR MET-11 IS THE INITIATOR.
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MEDLINE=98346338; PubMed=9682808;
Parker M.S., Larroque M.L., Campbell
 Cavia porcellus (Guinea pig).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
 PROSITE; PS00165; DEHYDRATASE_SER_THR; 1.
Threonine biosynthesis; Lyase; Pyridoxal phosphate.
BINDING 105 105 PYRIDOXAL PHOSPHATE (BY S. SEQUENCE 425 AA; 46682 MW; 2CC222DD201F09E0 CRC64;
 weneney K., Sutton G., Fitzhugh W., Fields C.A., Merrick J Scott J.D., Shirley R., Liu L.-I., Glodek A., Kelley J.M., Weidman J.F., Phillips C.A., Spriggs T., Hedblom E., Cotton M.D., Utterback T.R., Hanna M.C., Vguyen D.T., Saudek D.M., Brandon R.C. Gnehm C.D., Fritchman J.L., Fuhrmann J.L., Geodharan Venter T.
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 InterPro; IPR000634; -.
InterPro; IPR001926; -.
Pfam; PF00291; S_T_dehydratase; 1.
 EMBL; U32694; AAC21765.1; TIGR; HI0087; -.
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the European Bioinformatics Institute. There a
 -!- COFACTOR: PYRIDOXAL PHOSPHATE (I
 Science 269:496-512(1995).
 "Whole-genome random sequencing and assembly of Haemophilus influenzae \operatorname{Rd} ";
 STRAIN=RD / KW20 / MEDLINE=95350630;
 3 QVLGLQLPTP 12
 FUNCTION: BINDING MEDIATES SYNAPTIC
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 (RPL20A OR RPL18A2 OR YMR242C OR YM9408.04C) AND (RPL20B OR RPL18A1 OR RPL18B OR YOR312C OR O6116). Saccharomyces cerevisiae (Baker's yeast).
 EMBL; AF053328; AAC08993.1; -.
 SEQUENCE FROM N.A.
 PRINTS; PR01307; P2XRECEPTOR. PRINTS; PR01309; P2X2RECEPTOR.
 Pfam; PF00864; P2X_receptor; 1.
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 McDonnell J.M., Fushman D., Milliman C.L., Korsme "Solution structure of the proapoptotic molecule basis for apoptotic agonists and antagonists."; Cell 96:625-634(1999).
 EMBL; Z48756; CAA88652.1; ALT_SEQ.
EMBL; X90565; CAA62167.1; -.
EMBL; Z75220; CAA99632.1; -.
SGD; S0004855; RPL20A.
SGD; S0005839; RPL20B.
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 -!- SUBUNIT: FORMS HETERODIMERS EITHER WITH
BAX OR THE ANTI-APOPTOTIC PROTEIN BCL-2
-!- SUBCELLULAR LOCATION: PREDOMINANTLY CYTO
 Wang K., Yin X.-M., Chao D.T., Milliman C.L., "BID: a novel BH3 domain-only death agonist."; genes Dev. 10:2859-2869(1996).
 01-NOV-1997 (Rel. 35, Created)
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30-NAY-2000 (Rel. 39, Last annotation update)
BH3 INTERACTING DOMAIN DEATH AGONIST (BID).
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Mammalia; Eutheria; Rodentia;
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 -I- FUNCTION: INDUCES
EFFECT OF BCL-2.
 MEDLINE=99189748; PubMed=10089878;
 STRUCTURE BY NMR.
 MEDLINE=97078762; PubMed=8918887;
 NCBI_TaxID=10090;
 Mus musculus (Mouse).
 Ribosomal protein; Multigene :
SEQUENCE 174 AA; 20713 MW;
 InterPro; IPR002670; '-.
Pfam; PF01775; Ribosomal_L18ae;

 -!- MISCELLANEOUS: THERE ARE TWO GENES FOR L20 IN YEAST.
 -!- SIMILARITY: BELONGS TO THE L18AE FAMILY OF RIBOSOMAL PROTEINS.

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 Hypothetical protein. SEQUENCE 292 AA; 3
 PROSITE; PS00518; ZINC_FINGER_C3HC4; UNKNOWN_1.
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 EMBL; D79983; BAA11478.1;
 Nagase T., Seki N., Ishikawa K.-I., Tanaka A., Nomura N.; "Prediction of the coding sequences of unidentified human genes. The coding sequences of 40 new genes (KIAA0161-KIAA0200) deduced analysis of cDNA clones from human cell line KG-1."; DNA Res. 3:17-24(1996).
 121
 TISSUE=Bone marrow;
MEDLINE=96281124; PubMed=8724849;
 Eukaryota; Metazoa;
Mammalia; Eutheria;
 SEQUENCE FROM N.A.
 HYPOTHETICAL
 Homo sapiens (Human)
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Interpro; IPR000712;
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 Hartings H., Maddaloni M., Lazzarr Salamini F., Thompson R.D., "The OZ gene which regulates zein encodes a protein with structural activators."
 SEQUENCE FROM N.A.
STRALN-AC 1503 GM 1407;
MEDLINE-90016825; Pubmed-2798113;
Maddaloni M., di Fonzo N., Hartings H., La.
Salamini F., Thompson R.D., Motto M.;
"The sequence of the zein regulatory gene nucleic Acids Res. 17:7532-7532(1989).
 "The maize regulatory locus Opaque-2 encodes a DNA-binding protein which activates the transcription of the b-32 gene."; EMBO J. 10:617-624(1991).
 ЕМВО J. 8:2795-2801(1989).
 SEQUENCE FROM N.A.
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 MEDLINE=91160516; PubMed=2001677; Cohmer S., Maddaloni M., Motto M.
 MEDLINE=90059860; PubMed=2479535;
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 SIMILARITY: TO OTHER BZIP PROTEINS
 FUNCTION: INVOLVED IN THE REGULATION OF THE ENDOSPERM-SPECIFIC PRODUCTION OF ALBUMIN B-32 AND OTHER ZEIN PROTEINS. IT IS A TRANS-ACTING TRANSCRIPTIONAL ACTIVATOR THAT BINDS TO THE CONSENSUS SEQUENCE 5'-GATGAYRTGR-3'.
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 STRAIN-BRISTOL N2;
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 1- SIMILARITY: BELONGS TO THE NUCLEAR HORMONE RECEPTORS FAMILY.
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 -i- CATALYTIC ACTIVITY: ATP = 3',5'-CYCLIC AMP + PYROPHOSPHATE.
 Proc. Natl. Acad.
 "Two members of a widely expressed adenylyl cyclases.";
 Premont R.T.,
 MEDLINE=93028552; PubMed=1409703;
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 guanylate_cyc;
 GUANYLATE_CYCLASES_1;
GUANYLATE_CYCLASES_2;
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 & GAMMA SUBUNIT COMPLEX
ON: INTEGRAL MEMBRANE PRO
 Transmembrane;
 U.S.A. 89:9809-9813(1992).
 Ma H.-W.,
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M; 8B147E201C5DF601 C
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 membrane; Glycoprotein; Duplication CYTOPLASMIC (POTENTIAL).
 POLY-ALA.
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 Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Rattus
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 Ponnapalli M., ed subfamily of
 CALMODULIN-INSENSITIVE ADENYLYL
 (See http://www.isb-sib.ch/announce,
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Pfam; PF00211; guanylate_cyc; 2.

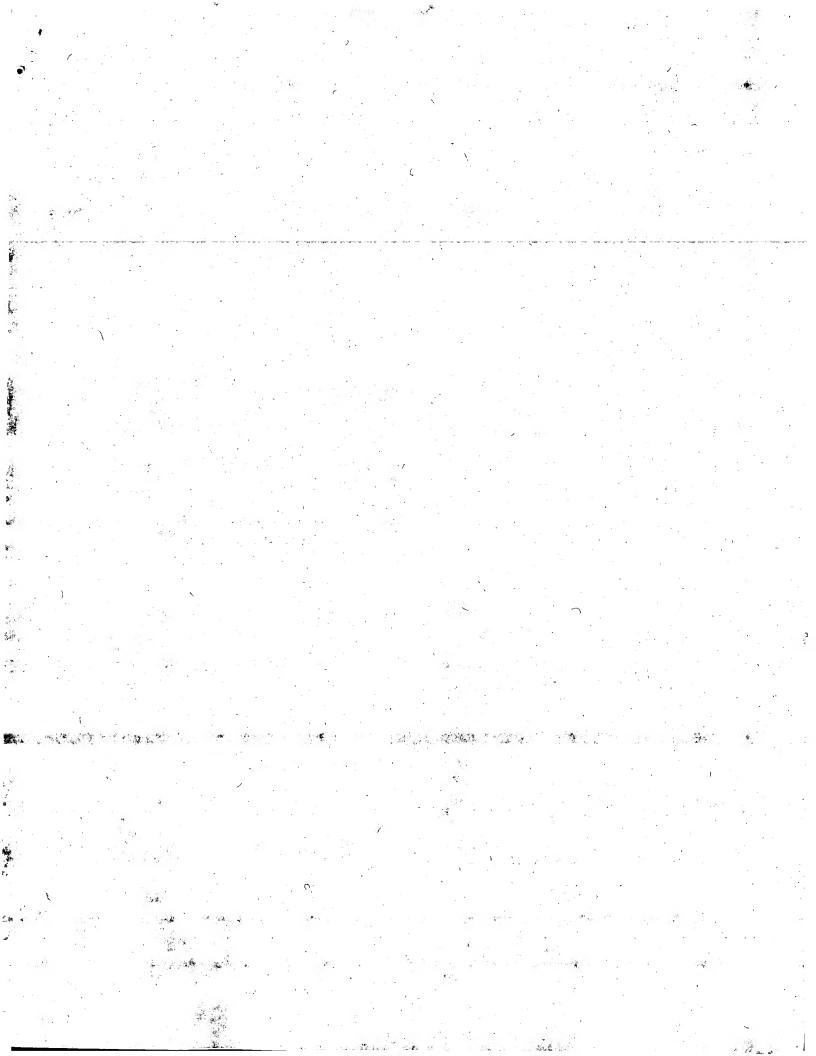
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PROSITE; PS50125; GUANYLATE_CYCLASES_2;
 use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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 PRANSMEM
 DOMAIN
 NIAMOC
 EMBL; Z29371; CAA82562.1;
HSSP; P19754; IAWK.
 TISSUE=Myocardium;
MEDLINE=94139935; PubMed=8307190;
Mellach J., Droste M., Kluxen F.W., Pfeuffer T.,
"Molecular cloning and expression of a novel type
 P40144;
01-FEB-1995
 RANSMEM
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 from rabbit myocardium.";
FEBS Lett. 338:257-263(1994)
 SEQUENCE FROM N.A., AND PARTIAL SEQUENCE
 ADCY5.
 01-FEB-1995 (Rel. 31, Last sequence update)
30 MAY-2000 (Rel. 39, Last annotation update)
ADENYLATE CYCLASE, TYPE V (EC 4.6.1.1) (ATP PYROPHOSPHATE-LYASE)
ADENYLYL CYCLASE).
 CYA5_RABIT
 CONCENTRATION RANGE. INTEGRAL MEMBRANE PROTEIN.
SUBCELLULAR LOCATION: INTEGRAL MEMBRANE PROTEIN.
TISSUE SPECIFICITY: MYCCARDIAL TISSUE.
DOMAIN: COMPOSED OF TWO HOMOLOGOUS DOMAINS.
PTM: THE N-TERMINUS IS BLOCKED.
SIMILARITY: BELONGS TO ADENYLYL CYCLASE CLASS-4/GUANYLYL CYCLASE
 CATALYTIC ACTIVITY: ATP = 3',5'-CYCLIC AMPENZYME REGULATION: INHIBITION BY CA(2+) IN
 FUNCTION: THIS A MEMBRANE BOUND, CA(2+)-INHIBITABLE ADENYLYL CYCLASE.
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 Transmembrane; Glycoprotein; Duplication.
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Maximum DB seq
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 Sequence:
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 Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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length: 2000000000
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 US-09-284-107-28
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Copyright (c) 1993 - 2000 Compugen Ltd.
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 4, 2001, 12:23:30; Search time 107.68 Seconds (without alignments) 13.062 Million cell updates/sec
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Q9rz01 deinococcus
Q9tyt0 caenorhabdi
 Q9nxb8 homo sapien
Q44982 caenorhabdi
 Q9pb61 xylella fas
070399 cavia porce
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|   | 58.3               | 58.3   | 58.3              | 60.0     | 60.0   | 60.0              | 60.0   | 60.0               | 60.0               | 60.0               | 60.0               | 60.0               | 60.0               | 60.0               | 60.0   | 60.0               | 60.0   | 60.0               | 60.0               | 60.0               | 60.0               | 60.0               | 60.0               | 60.0               | 60.0               | 60.0              |
|   | 145                | 115    | . 64              | 2151     | 1223   | 1048              | 639    | 609                | 568                | 492                | 437                | 419                | 408                | 395-               | 365    | 363                | 350    | 329                | 280                | 272                | 265                | 211                | 205                | 196                | 196                | 163               |
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|   |                    | `      |                   |          |        |                   |        |                    |                    | •                  |                    |                    |                    |                    |        |                    |        |                    |                    |                    |                    |                    |                    |                    |                    |                   |
|   | Q9wgf7 human papil |        | Q9vvg2 drosophila | trichomo | w      | Q9vyw5 drosophila |        | 09ug00 homo sapien | Q9nj97 caenorhabdi | Q9nj96 caenorhabdi | Q41835 zea mays (m | Q07795 sorghum bic | Q39532 coix lachry | Q9y3q8 homo sapien |        | Q9y7d0 aspergillus |        | 095144 homo sapien | Q9xet7 avena fatua | 095004 homo sapien | Q9s966 zea mays (m | Q9sfx8 arabidopsis | Q64763 avian adeno | Q9jk60 rattus norv | Q9jlt6 rattus norv | Q9u0w7 leishmania |

ALIGNMENTS

#### Q92663 Query Match Best Local Similarity Matches 12; Conserv Cells."; J. Clin. Invest. 90:2102-2109(1992). EMBL; L03418; AAA36049.1; HSSP; P12319; 1ALT. LWTERPRO; IPRO03006; LWTERPO; 179:30. PFAM; PF00047; 1g; 3. PRODOM; PD002534; -; 1 Q92663; Q92663; Q1-FEB-1997 Homo sapiens (Human). Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. SEQUENCE 01-FEB-1997 (TrEMBLrel. 02, 01-FEB-1997 (TrEMBLrel. 02, 01-OCT-2000 (TrEMBLrel. 15, Porges A.J., F Kimberly R.P.; MEDLINE=93055454; NCBI\_TaxID=9606; FC GAMMA RECEPTOR I. SEQUENCE FROM N.A. Novel Fc gamma receptor I family gene products in human mononuclear SSUE-BLOOD; PRELIMINARY; Redecha PubMed=1430234;. cha P.B., Doebele R., 42632 MW; 100.0%; Created) Last sequence update) Last annotation update) Score 60; DB 4; Pred. No. 0.0034; Mismatches 0 PRT; D33D59398CEEA699 CRC64; 374 Pan L.C., A A Length 374; Salmon J.E.

RESULT 2 Q92495 ID Q92495

PRELIMINARY;

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277 ELQVLGLQLPTP 288

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 Homo sapiens (Human).
Eukaryota; Metazoa; C
Mammalia; Eutheria; p
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 MEDLINE=93055454; PubMed=1430234; Porges A.J., Redecha P.B., Doebel Kimberly R.P.;
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 SEQUENCE
 PFAM; PF00047; ig; PRODOM; PD002534; -
 EMBL;
 Benech P.D., Sastry K., Iyer R.R., Eichbaum Q.G., Raveh D.P., Ezekowitz R.A., Ezekowitz R.A., "Definition of interferon gamma-response elements in a novel human Fc gamma_receptor gene (Fc gamma RIb) and characterization of the gene structure.
 278
 EMBL;
 EMBL; M91550;
 INTERPRO; IPR003006;
 structure
 SEQUENCE OF 1-224 FROM N.A. MEDLINE=93018827; PubMed=1402657;
 Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates;
 TISSUE=BLOOD;
 CD64 OR FC<GAMMA>RIB. Homo sapiens (Human).
 3zekowitz A.B.;
 SEQUENCE FROM N.A.
 NCBI_TaxID=9606;
 Submitted
 enech P.D., Sastry
 1 ELQVLGLQLPTP 12
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| 78 ELQVLGLRLPTP 28:
 S45704;
 P12319;
 S45707; AAD13842
S45708; AAD13842
 M91552;
 M91555; AAA58414
 1 Similarity 91.7
gamma receptor I family gene products in human mononuclear
 7 (TrEMBLrel.
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 PRELIMINARY;
 176:1115-1123(1992)
 AAD13842.
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 A84D464C70DD0F91 CRC64
 Craniata; Vertebrata; Euteleostomi;
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 PC burgers (Human).
Homo sapiens (Human).
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Yan Y., Li X., War
 Q9MZTO;
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 PRODOM;
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HSSP; P12319; 1ALT
 cells.";
 Q92637
 186 VKGLQLPTP 194
 'Novel Fc gamma receptor I family gene
 SEQUENCE FROM N.A.
 FC GAMMA RECEPTOR
 INTERPRO; IPRO03006; PFAM; PF00047; 1g; 1. PRODOM; PD002534; -;
 J. Clin. Invest. 90:210
EMBL; L03420; AAA35826
HSSP; P12319; 1ALT.
 SEQUENCE
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 94 VKGLQLPTP 102
 4 VLGLQLPTP 12
 Match
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 Craniata; V
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 Anopheles gambiae (African malaria mosquito).
Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;
Pterygota; Neoptera; Endopterygota; Diptera; Nematocera; Culiculae; Anopheles.
 01-MAY-2000 (TrEMBLrel. 13, 01-MAY-2000 (TrEMBLrel. 13, 01-CCT-2000 (TrEMBLrel. 15,
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Culicidae; Anopheles. NCBI_TaxID=7165;
 SERINE PROTEASE 22D.
 Q9NJS5
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 EMBL; AF105019; AAD54026.1; EMBL; AF250878; AAF69846.1;
 from Salmonella typhi
Nucleic Acids Res. 28
 Grotbeck
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MEDLINE-99296679; PubMed-10366528;
ROOKEY M.M., Sherbburne C., Lawley T.D., Taylor D.E.;
"Characterization of the Tra2 region of the IncHII p
 Salmonella
 Bacteria;
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 NCBI_TaxID=601;
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 beck E., Rose D.J., Taylor D.E.;
complete DNA sequence and analysis of
Salmonella typhi that is temperature s
elc Acids Res. 28:2177-2186(2000).
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 ELPVLGLQSTTP 288
 ELQVLGLQLPTP
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 41;226-239(1999).
 C.K., Lawley T.D.,
 Proteobacteria;
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 cloning and identification of full-length cDNA encoding ity Fc receptor for bovine IgG (Fc gamma RI)."; 1. Immunopathol. 75:151-159(2000).
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 Anopheles gambiae (African malaria mosquito).
Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda;
Pterygota; Neoptera; Endopterygota; Diptera; Nematoco
Culicidae; Anopheles.
 "Molecular characterization
SEQUENCE FROM Lennard N.;
 Caenorhabditis elegans
 01-OCT-2000
 "A modular chitin-binding prothemolymph in the mosquito Anop
Proc. Natl. Acad. Sci. U.S.A.
EMBL; AJ276428; CAB81934.1;
 Q9NATO
 Anopheles gambiae hemolymph.";
Insect Biochem. Mol. Biol. 30:35-46(2000).
EMBL; AF117751; AAD38337.3; -.
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 NCBI_TaxID=6239;
 Q9NAL2;
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inae; Caenorhabditis.
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 Endopterygota; Diptera; Nematocera;
 146811 MW;
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 protease associated wi
Anopheles gambiae.";
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 Lagueux M., Mueller
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 A Alvarenga R., Alves L.M.C., Araya J.E., Baia G.S., Baptista C.S.,
A Barros M.H., Bonaccorsi E.D., Bordin S., Bove J.M., Briones M.R.S.,
A Bueno M.R.P., Camargo A.A., Camargo L.E.A., Carraro D.M., Carrer H.,
Colauto N.B., Colombo C., Costa F.F., Costa M.C.R., Casta-Neto C.M.,
A Coutinho L.L., Cristofani M., Dias Neto E., Docena C., El-Dorry H.,
RA Facincani A.P., Ferreira A.J.S., Ferreira V.C.A., Ferro J.A.,
RA Fraga J.S., Pranca S.C., Franco M.C., Frohme M., Furlan L.R.,
A Fraga J.S., Pranca S.C., Franco M.C., Frohme M., Furlan L.R.,
RA Garnier M., Goldman G.H., Goldman M.H.S., Gomes S.L., Gruber A.,
RA Krieger J.E., Kuramae E.E., Laigret F., Lambais M.R., Leite L.C.C.,
RA Lemos E.G.M., Lemos M.V.F., Lopes S.A., Lopes C.R., Machado J.A.,
RA Machado M.A., Madeira A.M.B.N., Madeira H.M.F., Marino C.L.,
Martins E.M.E., Matsukuma A.Y.,
Monn D.H., Nagai M.A., Nascimento A.L.T., Monteiro-Vitorello C.B.,
RA Monn D.H., Nagai M.A., Nascimento A.L.T., Netto L.E.S.,
 В
 Qγ
 Query Match
Best Local (
 Matches
 Moon D.H., Nagai M.A., Nascimento A.L.T.O., Netto L.E.S.,
Nhani A. Jr., Nobrega F.G., Nunes L.R., Oliveira M.A.,
de Oliveira M.C., de Oliveira R.C., Palmieri D.A., Paris A.,
Peixoto B.R., Pereira G.A.G., Pereira H.A. Jr., Pesquero J.B.,
Quaggio R.B., Roberto P.G., Rodrigues V., de Rosa A.J.M.,
de Rosa V.E. Jr., de Sa R.G., Santelli R.V., Sawasaki H.E.,
da Silva A.C.R., da Silva A.M., da Silva F.R., Silva W.A. Jr.,
de Souza A.P., Ferenzi M.F., Truffi D., Tsai S.M., Tsuhako M.H.,
Vallada H., Van Sluys M.A., Verjovski-Almeida S., Vettore A.L.,
Zago M.A., Zatz M., Meidanis J., Setubal J.C.;
"maso M.A., Zatz M., Meidanis J., Setubal J.C.;
 Zago M.A., Zatz M., Meluanis J., The genome sequence of the plant pathogen Nature 406:151-157(2000).
 MEDLINE-20355717, Simpson A.J.G., Reinach F.C., Arruda r. Simpson A.J.G., Reinach F.C., Araya J.E., Alvarenga R., Alves L.M.C., Araya J.E., Bordin S.
 NTERPRO;
 NCBI_TaxID=2371;
 Bacteria;
 Xylella fastidiosa
 HYPOTHETICAL PROTEIN XF2283
 29PB61;
 29PB61
 investigating biology.";
Science 282:2012-2018(1998).
EMBL; ALI10477; CAB76742.1;
 222 KLOTLGIQLP 231
 "Genome sequence of the nematode
 10
 1 ELQVLGLQLP
 PRO; IPR001279;
PF00753; lactan
 Similarity 70.47; Conservative
 Proteobacteria;
 (TrEMBLrel.
 247 AA;
 (TrEMBLrel.
 lactamase_B;
 34299 MW;
 28826 MW;
 63.3%;
70.0%;
 15,
 gamma
 Last sequence update
Last annotation update
 Created)
 Score 38;
Pred. No.
 Arruda P.,
A99CE5F8A6B557B9
 subdivision;
 199EA394C41059B0 CRC64
 Mismatches
 annotation update)
 J.E., Baia G.S., B
 G
 .elegans: A
 317 AA.
 DB
 Xylella fastidiosa."
 Xanthomonas
 platform
 Length 247
 Indels
 Acencio
 for
 group;
 0;
 Gaps
 0
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Query Match

Score

38;

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070399
 Вb
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 Query Match
Best Local S
Matches 7
 Best Local
Matches
 EMBL; AK000342; BAA91097.1;
NON_TER 649 649
SEQUENCE 649 AA; 73060 My
 Q9NXB8 PRELIMINARY; PRT; 649 AA.
Q9NXB8;
01-OCT-2000 (TrEMBLrel. 15, Created)
01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-OCT-2000 (TrEMBLrel. 15, Last annotation update)
CDNA FLJ20335 FIS, CLONE HEP11429 (FRAGMENT).
 Eukaryota; Metazoa;
Mammalia; Eutheria;
NCBI_TaxID=9606;
 Submitted
 Kawakami T., Noguchi S., Itoh T., Sh
Nakajima Y., Mizuno T., Morinaga M.,
 Homo sapiens (Human)
 "NEDO human cDNA sequencing Submitted (FEB-2000) to the
 SEQUENCE
 PROSITE; PS01212; P2X_RECEPTOR; PRODOM; PD002383; -; 1.
Ionic channel; Transmembrane; I
 PRINTS; PR01307; P2XRECEPTOR.
PRINTS; PR01309; P2X2RECEPTOR.
 PFAM; PF00864; P2X_receptor;
 -!- SIMILARITY:
EMBL; AF053329;
 450
 INTERPRO;
 INTERPRO; IPRO01429;
 -!- SUBCELLULAR LOCATION: INTEGRAL MEMBRANE PROTEIN.
 Parker M.S., Larroque M
Hear. Res. 0:0;0(1998).
 SEQUENCE FROM N.A.
STRAIN-NIH STRAIN 2;
 Eukaryota; Metazoa;
 Cavia porcellus (Guinea pig)
 -! - SUBUNIT:
 NCBI_TaxID=10141;
 Mammalia; Eutheria;
 P2X2 RECEPTOR
 01-AUG-1998
01-OCT-2000
 01-AUG-1998
 070399;
 070399
 182 ELRIIGMDGPTP
 1 ELQVLGLQLPTP
 1 ELQVLGLQLPTP
 QAKLLGLQNPTP
 Similarity 7; Conserv
 FROM N.A.
 Similarity
6; Conser
 Shibahara T.,
 IPR003045;
 Conservative
 (TIEMBLrel. 07, Created)
(TIEMBLrel. 07, Last sequence update)
(TIEMBLrel. 15, Last annotation update)
 HOMO-
 Larroque M.L.,
 PRELIMINARY;
 Conservative
 SPLICE VARIANT
 Transmembrane; AA; 55771 MW;
 12
 461
 Chordata;
Primates;
 193
 12
 73060 MW;
 OR HETEROPOLYMERS (BY SIMILARITY)
 63.3%;
58.3%;
 Rodentia; Hystricognathi; Caviidae; Cavia.
 Chordata;
 TISSUE=ORGAN OF
 50.0%;
 Tanaka
 aka T., Nakamura Y
project.";
EMBL/GenBank/DDBJ
 Score 38; DB
Pred. No. 58;
3; Mismatches
 ω,
 Cambell J.M.,
 Catarrhini; Hominidae;
 Craniata; Vertebrata; Euteleostomi;
 P2X2-3.
 Ion transport; Rec
2B87E8B9F3CAC244
 Pred. No. 36;
4; Mismatches
18B9F87D373BC123 CRC64
 Craniata; Vertebrata;
 PRT;
 Shigeta K., Senba T.,
1., Ota T., Suzuki Y.,
 CORTI;
 501
 DB
58;
 B
 Bobbin
 11;
 Receptor
 databases
 2;
 Length 501;
 R.P.,
 Indels
 Euteleostomi;
 T.,
 Matsumura
Obayashi N
 Deininger P.;
 0;
 0;
 Gaps
 Gaps
 0,
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Query Match Best Local

63.3%;

4

Length 649;

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RESULT
084727
1D 08
AC 08
DT 01
DT 01
DT 01
DT 9H
GN PGI
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 Query Match
Best Local Similarity
Matches 7; Conserv
 Matches
 SEQUENCE FROM N.A. STRAIN-BRISTOL N2; Waterston R.;
 Wilson R., Ainscough R., Anderson K., Baynes C., Berks M.,
Bonfield J., Burton J., Connell M., Copsey T., Cooper J., Coulson A.,
Craxton M., Dear S., Du Z., Durbin R., Favello A., Fulton L.,
Gardner A., Green P., Hawkins T., Hillier L., Jier M., Johnston L.,
Jones M., Kershaw J., Kirsten J., Laister N., Latreille P.,
Jones M., Kershaw J., Kirsten J., Laister N., Latreille P.,
Lightning J., Lloyd C., Mcmurray A., Mortimore B., O'Callaghan M.,
Parsons J., Percy C., Rifken L., Roopra A., Sulston J.,
Smaldon N., Smith A., Sonnhammer E., Staden R., Sulston J.,
Thierry-Mieg J., Thomas K., Vaudin M., Vaughan K., Waterston R.,
Watson A., Weinstock L., Wilkinson Sproat J., Wohldman P.;
 01-NOV-1998
01-OCT-2000
 084727;
01-NOV-1998
 084727
 STRAIN=BRISTOL N2;
Tin-Wollam A., Graves T.,
 01-JUN-1998
01-NOV-1998
 044982
 SEQUENCE FROM N.A.
STRAIN-BRISTOL N2;
 Eukaryota; Metazoa;
Rhabditidae; Pelode
 C54G6.4
 C54G6.4 PROTEIN
 044982;
Chlamydia trachomatis.
 PHOSPHOGLYCERATE
 SEQUENCE
 NON_TER
 EMBL; AF043698; AAB97562.1;
 Submitted
 Submitted
 SEQUENCE FROM N.A.
 Caenorhabditis elegans
 01-JUN-1998
 Nature 368:32-38(1994).
 MEDLINE=94150718; PubMed=7906398;
 86
 66 LQIAGLSLPLP 76
 14
 13
 2 LQVLGLQLPTP 12
 3 QVLGLQLPTP
 2 Mb of
 QQLGVQLPVP 107
 Similarity 7; Conserv
 (JAN-1998)
 (JAN-1998)
 134 AA;
 contiguous nucleotide sequence
 3 (TrEMBLrel. 06,
8 (TrEMBLrel. 06,
1 (TrEMBLrel. 08,
 (TrEMBLrel. 08, (TrEMBLrel. 15,
 Conservative
 PRELIMINARY;
 Conservative
 PRELIMINARY;
 Peloderinae;
 12
 (FRAGMENT).
 MUTASE
 13720 MW;
 Nematoda; Chromadorea; Cinae; Caenorhabditis.
 61.7%;
70.0%;
 ξ
 to the EMBL/GenBank/DDBJ
 the EMBL/GenBank/DDBJ
 Ozersky P.;
 Created)
Last sequence update)
Last annotation update)
 Last sequence update)
Last annotation update)
 Score 38; DB Pred. No. 76; 1; Mismatches
 ۲
 Created)
 1;
 Score 37;
Pred. No.
 PRT;
 PRT;
 24F44A10FA48CDEE CRC64;
 Mismatches
 226
 134
 DВ
 A
 from chromosome III of C.
 5.
 Rhabditida;
 databases
 databases
 2
 Length 134;
 Indels
 Indels
 Rhabditoidea;
 0
 0
 Gaps
 Gaps
 R.
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 Matches
 Query Match
Best Local
 Matches
 Query Match
Best Local
 White O., Eisen J.A., Heidelberg J.F., Hickey E.K., Dodson R.J., Haft D.H., Gwinn M.L., Nelson W.C., Ric Moffat K.S., Qin H., Jiang L., Pamphile W., Crosby M. Vamathevan J.J., Lam P., McDonald L., Utterback T.,
 01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-UUN-2000 (TrEMBLrel. 14, Last annotation update)
 Q9RZ01;
 Q9RZ01
 PFAM; PF00300; PGAM; 2.
PROSITE; PS00175; PG_MUTASE; UNKNOWN_1.
SEQUENCE 226 AA; 25785 MW; 82AE9C780D400B8D CRC64;
 Chlamydia trachomatis.";
Science 282:754-759(1998).
EMBL; AE001343; AAC68317.1;
 STRAIN=D/UW-3/CX;
MEDLINE=99000809; PubMed=9784136;
 SEQUENCE
 PFAM; PF01614; SEQUENCE 253
 EMBL;
 radiodurans R1.";
Science 286:1571-1577(1999).
 Makarova K.S., Aravind L.,
Ketchum K.A., Nelson K.E.,
 SEQUENCE FROM N.A.
 NCBI_TaxID=1299;
 Bacteria; Thermus/Deinococcus
 Deinococcus radiodurans
 DRA0152.
 TRANSCRIPTIONAL REGULATOR,
 Davis R.W.;
 Mitchell W.P., Olinger L.,
 SEQUENCE FROM
 NCBI_TaxID=81
 Bacteria; Chlamydiales; Chlamydiaceae; Chlamydia
 TIGR;
 Genome Sequence of the Radioresistant Bacterium Deinococcus
 Fraser C.M.;
 MEDLINE=20036896; PubMed=10567266;
 [NTERPRO;
 HSSP;
 INTERPRO; IPR000285; -.
 Genome sequence of an obligate
205
 N
LAALGVSLPTP
 LQVLGLQLPTP 12
 AE001862; AAF12212.
 P00950; 5PGM
 DRA0152;
 7; Conserv
 Similarity
8; Conserv
 IPR001345; -.
 Conservative
 PRELIMINARY;
 Conservative
 N.A.
 IclR;
AA;
 205
215
 11
 27299
 61.7%;
 Lammel C.J., Fa, Tatusov R.L.,
 MW;
 cDonald I., Utterback T., Zalewski C.,
Daly M.J., Minton K.W., Fleischmann R.D.,
Salzberg S., Smith H.O., Venter J.C.,
 ICLR FAMILY.
 ۲;
 Score 37;
Pred. No.
 Score 37; DB Pred. No. 39;
 group;
 PRT;
 DBB8F05ABE6F65A8 CRC64;
 Mismatches
 Mismatches
 intracellular pathogen of humans:
 Deinococcales; Deinococcus
 253
 DB 44;
 Fan J.,
.., Zhao (
 DB
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 2
 2;
 Crosby M., Shen M., back T., Zalewski C.,
 Q., Koonin E.V.,
 Ψ
 2
 Length 226
 Length 253;
 Richardson D.L.,
 Indels
 Indels
 Peterson J.D.,
 0;
 0;
 Gaps
 Gaps
 .
 0,
 0
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Search completed: June

4

2001, 12:23:31

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GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.

OM protein - protein search, using sw model

Run on:
June 4, 2001, 12:13:33; Search time 106.14 Seconds (without alignments)
6.463 Million cell updates/sec

Title:
US-09-284-107-29
Perfect score: 68
```

Perfect score: 68
Sequence: 1 VWFHVLFYLAVG 12
Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5
Searched: 390729 seqs, 57163235 residues

Total number of hits satisfying chosen parameters: 390729

Minimum DB seq length: 0 Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database: A\_Geneseq\_0401:\*

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2: /SIDS6/gcgdata/geneseq/geneseqp/AA1981.DAT:\*

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10: /SIDS6/gcgdata/geneseq/geneseqp/AA1989.DAT:\*

11: /SIDS6/gcgdata/geneseq/geneseqp/AA1990.DAT:\*

12: /SIDS6/gcgdata/geneseq/geneseqp/AA1993.DAT:\*

13: /SIDS6/gcgdata/geneseq/geneseqp/AA1993.DAT:\*

14: /SIDS6/gcgdata/geneseq/geneseqp/AA1993.DAT:\*

15: /SIDS6/gcgdata/geneseq/geneseqp/AA1993.DAT:\*

17: /SIDS6/gcgdata/geneseq/geneseqp/AA1995.DAT:\*

18: /SIDS6/gcgdata/geneseq/geneseqp/AA1997.DAT:\*

19: /SIDS6/gcgdata/geneseq/geneseqp/AA1999.DAT:\*

20: /SIDS6/gcgdata/geneseq/geneseqp/AA1999.DAT:\*

21: /SIDS6/gcgdata/geneseq/geneseqp/AA1999.DAT:\*

22: /SIDS6/gcgdata/geneseq/geneseqp/AA1999.DAT:\*

23: /SIDS6/gcgdata/geneseq/geneseqp/AA1999.DAT:\*

24: /SIDS6/gcgdata/geneseq/geneseqp/AA1999.DAT:\*

25: /SIDS6/gcgdata/geneseq/geneseqp/AA1999.DAT:\*

26: /SIDS6/gcgdata/geneseq/geneseqp/AA1999.DAT:\*

27: /SIDS6/gcgdata/geneseq/geneseqp/AA1999.DAT:\*

28: /SIDS6/gcgdata/geneseq/geneseqp/AA1999.DAT:\*

29: /SIDS6/gcgdata/geneseq/geneseqp/AA1999.DAT:\*

20: /SIDS6/gcgdata/geneseq/geneseqp/AA1999.DAT:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

# SUMMARIES

|   |              |             |             |             |             |        |            |              |              |                  |                   |             | , |
|---|--------------|-------------|-------------|-------------|-------------|--------|------------|--------------|--------------|------------------|-------------------|-------------|---|
|   | 11           | 10          | 9           | 60          | . 7         | σ      | U          | 4            | ω            | Ν                | _                 | No.         |   |
|   | 68           | 83          | 89          | 83          | 89          | 83     | 83         | 68           | 68           | 68               | 89                | Score       |   |
|   | 100.0        | 100.0       | 100.0       | 100.0       | 100.0       | 100.0  | 100.0      | 100.0        | 100.0        | 100.0            | 100.0             | Match 1     |   |
| • | 399          | 374         | . 374       | 374         | 374         | 374    | 374        | 374          | 344          | 344              | 12                | Length DB   |   |
|   | 21           | 21          | 21          | 19          | 19          | 19     | 17         | 17           | 21           | 17               | 19                | DB          |   |
|   | B43683       | Y96226      | Y96134      | W97834      | W97833      | W80448 | W00859     | R91438       | Y96183       | R91439           | W60561            | 븅           |   |
|   |              |             |             |             |             |        |            |              |              |                  |                   |             |   |
|   |              |             |             |             |             |        |            |              |              |                  |                   |             |   |
|   |              |             |             |             |             |        |            |              |              |                  |                   | :           |   |
|   | -            | Human       | Human       | Human       | Human       |        | Human      | Human        | Human        | Human            | Oligo             | Description |   |
|   | cancer assoc | high affini | macrophage- | Fc receptor | Fc receptor | w      | FCRI (CDNA | FCRI (CDNA o | macrophage-s | Human FcRI (cDNA | Oligopeptide from | otion       |   |
|   | (3           | ~           | J)          |             |             |        | ()         | á            | O1           | ()               |                   | I           |   |

Identifying peptide(s) binding specifically to protein target - by expressing on phage surface and testing for binding to immobilised oligo; peptide derived from the target, useful for, e.g. identifying specific antibodies

WPI; 1998-240964/21.

De Kruif CA,

Logtenberg T;

(UYUT-) RIJKSUNIV UTRECHT.

08-OCT-1996;

96EP-0202791.

| 45     | 44     | 43                 | 42     | 41     | 40     | 39     | 38     | 37     | 36     | 35                 | 34                 | <u>ω</u> | 32            | 31                 | 30     | 29     | 28     | 27                 | 26     | 25     | 24                 | 23                 | 22                 | 21                 | 20                 | 19                 | 18           | 17           | 16     | 15     | 14                 | 13     | 12                 |
|--------|--------|--------------------|--------|--------|--------|--------|--------|--------|--------|--------------------|--------------------|----------|---------------|--------------------|--------|--------|--------|--------------------|--------|--------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------|--------------|--------|--------|--------------------|--------|--------------------|
| 38     | 38     | 3 <b>8</b> .       | 38     | 3<br>8 | 38     | 38     | 38     | 38     | 38     | 38                 | 38                 | 38       | <u>ن</u><br>8 | 38                 | 38     | 38     | 38     | 38                 | 38     | 38     | 38                 | 38                 | 38                 | 38                 | 38.5               | 42                 | 42           | 42           | 49     | 49     | 49                 | 51     | 64                 |
| 5      | 5.     | 5                  | 5      | 5      | 5      | 5      | 5      | 5      | 5      | 5                  | 5                  | 5        | 5.            | 5                  | 5      | 5      | 5      | 5                  | 5      | 5      | 5                  | 5                  | 5.                 | 5                  | ٠,                 | -                  | ۲.           | 61.8         | Ņ      | 2      | 2                  | 75.0   | 94.1               |
| 1176   | 610    | 610                | 423    | 423    | 398    | 398    | 398    | 395    | 391    | 391                | 327                | 327      | 281           | 273                | 273    | 267    | 267    | 267                | 266    | 266    | 266                | 203                | 203                | 65                 | 1084               | 1.64               | 98           | 96           | 373    | 257    | 243                | 377    | 410                |
|        |        |                    |        |        |        |        |        |        |        |                    |                    |          |               |                    |        |        |        |                    |        |        |                    |                    |                    |                    |                    |                    |              |              | 21     | 21     | 21                 | 20     | 12                 |
| G49881 | W38417 | W43410             | G42851 | G09391 | G50427 | G05034 | Y24479 | Y24478 | G42852 | G09392             | G50428             | · G05035 | G14199        | G37548             | G06464 | G50429 | G14200 | G05036             | G37549 | G06465 | Y16106             | G37550             | G06466             | G33208             | B42371             | G40829             | G40830       | G40831       | G50154 | G50155 | G50156             | W86195 | R12428             |
|        |        |                    |        |        |        |        |        |        |        |                    |                    |          |               |                    |        |        |        |                    |        | ٠      |                    |                    |                    |                    |                    |                    |              |              |        |        |                    |        |                    |
|        |        |                    |        |        |        |        |        |        |        |                    |                    |          |               |                    |        |        |        |                    |        |        |                    |                    |                    |                    |                    |                    |              |              |        | -      |                    | :      |                    |
| <br>Q. | 4      | Yeast acylcoenzyme | is     | ŝ      | is tha | -      |        |        | ۳.     | Arabidopsis thalia | Arabidopsis thalia | is       | is            | Arabidopsis thalia | is tha | is tha | is tha | Arabidopsis thalia | Š      | S      | A formate transpor | Arabidopsis thalia | Arabidopsis thalia | Zea mays protein f | Human ORFX ORF2135 | Zea mays protein f | mays protein | mays protein |        | İS     | Arabidopsis thalia | н.     | Hybrid Fc(gamma)RI |

## ALIGNMENTS

| 7      | 3                                                           |
|--------|-------------------------------------------------------------|
| W60561 | - 1                                                         |
|        | W60561 standard; peptide; 12 AA.                            |
|        |                                                             |
|        | W60561;                                                     |
|        |                                                             |
|        | 18-AUG-1998 (first entry)                                   |
| ×      |                                                             |
|        | Oligopeptide from extracellular domain of CD64.             |
| X      |                                                             |
|        | Extracellular domain; CD64; identification; antibody;       |
|        | immunohistochemical; immunofluorescent analysis; detection; |
|        | cell transformation; mutation; anti; oncogene.              |
| X      | •                                                           |
|        | Synthetic.                                                  |
|        |                                                             |
|        | WO9815833-A1.                                               |
| ×      |                                                             |
|        | 16-APR-1998.                                                |
| ×      |                                                             |
| PF (   | 07-OCT-1997; 97WO-NL00557.                                  |
|        |                                                             |

CDNA

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PR PR XX
 ET XX
 В
 QΥ
 Matches
 Query Match
Best Local 9
 13-JUL-1989;
13-JUL-1990;
18-OCT-1993;
WPI; 1996-200279/20
N-PSDB; T14718.
 Aruffo A,
 01-DEC-1992;
25-FEB-1988;
 (GEHO) GEN HOSPITAL CORP
 25-FEB-1988;
 09-APR-1996
 Misc-difference
 Cell surface antigen; cloning; immunoselection; immunotherapy; therapy; diagnosis; vector; FCRI; FC receptor.
 surface of a replicable display package, synthesising oligopeptides derived from the target protein on a solid phase, and testing for binding between the peptide and oligopeptides. The method is used to screen large that bind to cell markers or that can differentiate between different froms of the same protein, including bispecific antibodies that bind to epitopes on the same protein, including bispecific antibodies that bind to epitopes on different molecules. The genes/oligonucleotides that encode selected peptides can be isolated and used for recombinant production of immunofluorescent analysis, and also to detect cell transformation caused by manufactured in contractions.
 US5506126-A.
 Homo sapiens
 Human FcRI (cDNA clone p98 product).
 30-OCT-1996
 R91439
 R91439
 Sequence
 by mutation in (anti)oncogenes.
 affinity binding to the CD64-covered rods. The specification describes a method for the identification of a peptide able to bind specifically to a target protein. The method comprises displaying the peptide on the surface of a replicable display package, synthesising oligopeptides
 adjacent, non-overlapping sets. Peptide specific phages were affinity binding to the CD64-covered rods. The specification
 Synthetic oligopeptides W60537-61 are domain of CD64. They were synthesised
 Example 1; Page 29; 40pp; English
 Local Similarity 100 hes 12; Conservative
 1 VWFHVLFYLAVG 12
||||||||||||
1 vwfhvlfylavg 12
 standard; Protein; 344
 Seed B;
 12 AA;
 (first entry)
 92US-0983647.
88US-0160416.
89US-0379076.
90US-0553759.
 88US-0160416
 Location/Qualifiers 58
 /note= "amino acid
translated
 100.0%;
 0;
 AA.
 Score 68; DB 19;
Pred. No. 1.1e-05;
 Mismatches
 58 is Leu
product"
 58
 derived from the extracellular on 25 polyethylene rods as
 in clone
 Length 12;
 Indels
 p135
 0;
 Gaps
```

```
RESULT
Y96183
 Дδ
 Qy
 CCCCCCXXXX PT
 Query Match
Best Local Similarity
Matches 12; Conserv
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc:difference
 Misc-difference
 Misc~difference
 Misc-difference
 panning; immunodiagnosis; diagnosis; immunotherapy; gene therapy; immune disorder; infection; asthma; immune-complex disease; amyloidosis; multiple scierosis; parasitic disease; Fc receptor.
 Misc-difference
 The amino acid sequence (R91439) of human FCRI was detd. from a cDNA clone, p98 (T14718), obtd. from a cDNA library using an immunoselection cloning method. FCRI is a high affinity receptor for the Fc portion of IgG, normally located on cell surfaces of macrophages. 2 Other cDNA clones, p135 (T14717) and p90 (T14719), of the p98 product is truncated compared with those of the p135 and p90 products (see also R91438 and W00859). A fusion protein of FCRI and a receptor ligand will be helpful in increasing the potency of antibodies in therapy.
 Homo sapiens
 Macrophage; FCRI; cell surface antigen; human; immunoselection;
 Human macrophage-specific FcRI.
 19-DEC-2000 (first entry)
 Y96183 standard; Protein; 344 AA
 Sequence
 Cloning of cDN
of diagnostic
 1 VWFHVLFYLAVG 12
||||||||||
|289 vwfhvlfylavg 300
 Example 10; Column 55-56; 79pp;
 344 AA;
 Conservative,
 /note=
 /note=
? 51
 /note=
123
 /note=
122
 /note=
117
 NA encoding cell surface antigen - useful for isolation and therapeutic proteins
 116
 /note-
64
 /note=
60
 /note=
 /note=
55
 /note=
50
 38
 Location/Qualifiers
 note=
 'note=
 100.0%;
 "encoded by
 "encoded by
 "encoded by
 "encoded by CAG"
 "encoded
 encoded by
 "encoded
 "encoded by TCC"
 "encoded
 "encoded
 "encoded by CGG"
 "encoded by ACC"
 "encoded by
 0;
 Score 68; DB 1/;
Pred. No. 0.00038;
 уd
 γď
 by ccc"
 Ьy
 English
 ACC"
 TTC"
 GCA"
 GAA"
 CAA"
 CAC "
 ACC"
 Length 344;
 Indels
 0;
 Gaps
```

0;

Misc-difference

0

```
XX
 01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
23-MAR-1990;
13-JUL-1990;
 based upon transient expression of a CSA in eukaryotic cells and physical selection of cells expressing the antigen by adhesion to (panning on) an antibody-coated substrate such as a culture dish. CSA nucleic acids isolated by the method of the invention, and the proteins they encode, are useful for immunodiagnostic and immunotherapeutic applications, including the diagnosis and
 ability to
 immunotherapeutic applications, including the diagnosis and treatment of immune-mediated infections, diseases, and disorders in animals, including humans. These disorders include asthma,
 The present sequence is that of a human macrophage specific FcRI, as deduced from cDNA clone pl35 (see A50631), which was isolated from a cDNA library expressed in COS cells using a novel method of the invention designed to isolate CSA nucleic acids. The method is
 Example 10;
 Isolated nucleic acid molecule encoding the CD19 cell surface antigen, useful for immunodiagnosis and immunotherapy of immune-mediated infections or disorders, e.g. asthma, immune-complex disease, parasiti
 Stamenkovic I,
 28-OCT-1998;
 29-AUG-2000
 US6111093-A
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 (GEHO) GEN HOSPITAL CORP
 immune-complex disease, amyloidosis, parasitic diseases or multiple sclerosis. FCRI is a high affinity receptor for the Fc portion of
 2000-586382/55
)B; A50631.
 normally located on the cell surfaces of macrophages.
 Column
 900S-0498809
900S-0553759
 88US-0160416
89US-0379076
 92US-0983647
 98US-0181612
 /note=
333
 /note=
332
 /note=
 /note=
213
 /note=
139
 /note=
134
 Seed
 305
 268
 220
 136
 /note=
129
 /note-
 /note=
 /note=
 /note=
 /note=
 /note=
 note-
 'note=
 53-55; 75pp;
 "encoded
 "encoded by
 "encoded by
 "encoded by
 "encoded by CCT"
 "encoded by
 "encoded by
 "encoded by
 "encoded by
 "encoded
 "encoded
 "encoded by
 "encoded by
 "encoded by
 "encoded
 bonding, or
 Ьy
 bу
 ру
 Ьy
 English
 AAT
 TTC"
 CCA"
 GTT "
 AAT "
 CAT"
 GGT
 AAC"
 GIG"
 CGT"
 CAA"
 AAT"
 GGC
 immune-complex disease, parasitic
 cause it to occur on
```

The

Sequence sequence. be helpful

374

in increasing the potency of antibodies in

```
8 x 0 0 0 0 0 0 0 0 0 x &
 RESULT
R91438
 밁
 δÃ
 Query Match
Best Local Similarity
Taches 12; Conserve
 SSSS
 13-JUL-1989;
13-JUL-1990;
18-OCT-1993;
 Cloning of cDNA encoding cell surface antigen - useful for isolation of diagnostic and therapeutic proteins % \left(1\right) =\left\{ 1\right\}
 Sequence
 protein of FCRI and a receptor ligand will be helpful to increase the potencies of antibodies in therapy.
 for the Fc portion of IGG, normally located on cell surfaces of macrophages. Another isolated cDNA clone, p90 (T14719), coded for a variant (W00859) of the FcRI sequence, and a third clone, p98 (T14718), coded for an FcRI (R91439) having a different C-terminal sequence. A fusion protein of FcRI and a receptor ligand will
 The amino acid sequence (R91438) of human FCRI was detd. from cDNA clone, p135 (T14717), obtd. from a cDNA library using an
 09-APR-1996
 Misc-difference
 Cell surface antigen; cloning;
 Human FcRI (cDNA clone pl35 product).
 R91438;
 R91438 standard; Protein; 374 AA
 WPI; 1996-200279/20
N-PSDB; T14717.
 Aruffo'A,
 (GEHO) GEN HOSPITAL CORP
 01-DEC-1992;
25-FEB-1988;
 25-FEB-1988;
 US5506126-A
 Homo sapiens
 30-OCT-1996
 Example 10; Column 55-56; 79pp; English.
 Misc-difference
 therapy; diagnosis;
 1 WWFHVLFYLAVG 12
|||||||||||
| 289 vwfhvlfylavg 300
 mmunoselection cloning method.
 4
 Seed B;
 344 AA;
 Conservative
 (first entry)
 92US-0983647.
88US-0160416.
89US-0379076.
90US-0553759.
93US-0139273.
 88US-0160416
 Location/Qualifiers
 58
 /note=
 /note=
 vector; FcRI; Fc receptor
 100.0%;
 "amino acid
translated
 "amino acid
 translated
 obtd.
 9
 Score 68;
Pred. No.
 immunoselection;
 Mismatches
 FcRI is a high affinity receptor
 25 is Thr
products"
 58 is val in product"
 0.00038;
 DB
 21;
 in
 0,
 p90 clone
 clone p90 and
 immunotherapy;
 Length 344;
 Indels
 0;
 998
 Gaps,
```

Query Match Best Local

Length 374;

```
밁
 Š
 Qy
 В
 Query Match
Best Local Similarity
Matches 12; Conserv
 Matches
 01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
18-OCT-1993;
 cDNA clone, p90 (714719), obtd. from a cDNA library using an immunoselection cloning method. FCRI is a high affinity receptor for the FC portion of IgG, normally located on cell surfaces of macrophages. Other cDNA clones (see also T14717-18) coded for variants (R91438-39) of the FCRI sequence. A fusion protein of FCRI and a receptor ligand will be helpful in increasing the potency of antibodies in therapy.
 The amino acid sequence (W00859) of human FCRI was detd. CDNA clone, p90 (T14719), obtd. from a cDNA library using
 Cloning of cDNA encoding cell surface antigen - useful for isolation of diagnostic and therapeutic proteins % \left(1\right) =\left\{ 1\right\}
 Sequence
 WPI; 1996-200279/20.
N-PSDB; T14719.
 Aruffo
 289
 Example 10; Column 55-56; 79pp; English.
 (GEHO) GEN HOSPITAL CORP
 25-FEB-1988;
 09-APR-1996
 US5506126-A.
 Homo sapiens
 Cell surface antigen; cloning; immunoselection; therapy; diagnosis; vector; FCRI; FC receptor.
 Human FcRI (cDNA clone p90 product).
 30-OCT-1996
 W00859;
 W00859 standard;
 1 VWFHVLFYLAVG
||||||||||
| 289 vwfhvlfylavg
vwfhvlfylavg 300
 VWFHVLFYLAVG 12
 S
 A,
 ch 100.0%; Score 68; DB 17; Similarity 100.0%; Pred. No. 0.00042; 12; Conservative 0; Mismatches 0;
 Seed
 374 AA;
 Conservative
 (first
 88US-0160416.
89US-0379076.
90US-0553759.
93US-0139273.
 92US-0983647
 88US-0160416
 Protein;
 300
 12
 100.0%;
 374
 0
 Score 68; DB 17;
Pred. No. 0.00042;
 A
 Mismatches
 0;
 immunotherapy;
 Length 374;
 Indels
 Indels
 receptor
 0;
 0;
 Gaps
 Gaps
 0,
 0;
```

RESULT W80448

This is the amino acid sequence of human Fc receptor I (FCRI), and deduced from cDNA clone pi35 (see V63456) isolated using a rapid immunoselection cloning method from a cDNA library expressed in cells. The cDNA library was constructed from polyA RNA of cells from a single patient undergoing extracorporeal interleukin-2

cos

Example 10;

Column 53-54; 75pp; English

```
01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
21-MAY-1997;
 New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially human lymphocyte antigenic sequences
 WPI; 1998-609251/51
N-PSDB; V63456.
 Aruffo A,
 21-MAY-1997;
 US5830731-A
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 (GEHO) GEN
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 cloning.
 Homo sapiens
 Fc receptor I; FcRI;
 Human Fc
 07-JUN-1999
 W80448 standard; Protein; 374 AA.
 Seed B;
 HOSPITAL CORP
 (first entry)
 92US-0983647.
88US-0160416.
89US-0379076.
90US-0553759.
 /note=
- 85
 /note=
171
 97US-0861205
 97us-0861205
 /note=
159
 /note=
 141
 256
 60
 Location/Qualifiers
 /note=
 /note=
 note=
 /note=
 'note=
 note= "encoded by CTG"
 note=
 note= "encoded by CTC"
 note= "encoded by TGG"
 cell surface
 "encoded
 "encoded by
 "encoded by GGG"
 "encoded
 "encoded
 "encoded by
 "encoded by
 "encoded by AAT"
 "encoded
 "encoded by
 "encoded
 Ьy
 þу
 Ьy
 bу
 by GAC"
 · GTC"
 CCC.
 antigen; lymphocyte;
 AAC"
 "DBB
 CAA"
 TCC"
 ATG"
 CCC"
 human;
```

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밁
 Š
 Query Match
Best Local Similarity
Matches 12; Conserv
 adhesion to an antibody-coated substrate. The method is useful for the isolation and cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V63442-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated genes can be expressed in a prokaryotic or enkaryotic host cells to produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V63444) which allow the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and
 Misc-difference 141
 Fc receptor I;
cloning.
 Human Fc receptor I
Misc-difference
 Misc-difference
 Misc-difference 103
 Misc-difference
 07-JUN-1999
 W97833;
 immunotherapeutic applications, treatment of immune-mediated in
 were also obtained. A novel method for cloning cDNAs from mammalian expression libraries is based on transient expression of an antigen by in eukaryotic cells and selection of cells expressing the antigen by
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Homo sapiens
 W97833 standard; Protein; 374 AA.
 Sequence
 induction therapy.
 1 VWFHVLFYLAVG 12
 374 AA;
 Conservative
 (first entry)
 immune-mediated infections, diseases, and disorders of
 /note=
3 85
 /note-
77
 FCRI;
 /note=
171
 /note-
159
 /note=
99
 /note=
 /note=
45
 /note-
176
 Location/Qualifiers
 /note-
 /note= "encoded by CTC"
 /note=
 'note= "encoded by TGG"
 100.0%; Sc.
100.0%; Pr
 Clones p90 (see X07372) and p98 (s
A novel method for cloning cDNAs
 cell surface antigen; lymphocyte; human;
 "encoded by CAA'
 "encoded by
 "encoded by
 "encoded by
 "encoded
 "encoded by
 "encoded by
 "encoded by
 "encoded by AAT"
 "encoded
 "encoded by GAC"
 Score 68; DB 19;
Pred. No. 0.00042;
 bу
 bу
 Mismatches
 including the diagnosis and
 GGC
 CCC
 CTG"
 GTC'
 ATG"
 AAC"
 CCC.
 TCC
 0
 Length 374;
 Indels
 (see X07373)
 0;
 Gaps
 0;
```

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QΥ
 TXXX
 밁
 RESULT
 XX
 Query Match
Best Local Similarity
Matches 12; Conser
 01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
21-MAY-1997;
 has been used to clone genes (see V63442-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated genes can be expressed in a prokaryotic or eukaryotic host cells to produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V63444) which allow the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and immunotherapeutic applications, including the diagnosis and treatment of immune-mediated infections, diseases, and disorders of
Human Fc receptor I.
 07-JUN-1999 (first entry)
 W97834 standard;
 expression libraries is based on transient expression of an antigen in eukaryotic cells and selection of cells expressing the antigen by adhesion to an antibody-coated substrate. The method is useful for
 W97834;
 Sequence
 adhesion to an antibody-coated substrate. The method is useful the isolation and cloning of any protein which can be expressed transported to the cell surface membrane of a eukaryotic cell.
 Example 10; Column 53-54; 75pp; English
 New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially
 WPI; 1998-609251/51.
N-PSDB; X07372.
 Aruffo A,
 21-MAY-1997;
 03-NOV-1998
 were also obtained.
 from a single patient undergoing extracorporeal interleukin-2 induction therapy. Clones pl35 (see V63456) and p98 (see X07373)
 cells.
 human lymphocyte antigenic sequences
 (GEHO) GEN HOSPITAL CORP
 US5830731-A
 immunoselection cloning method from a cDNA library expressed
 289 vwfhvlfylavg
 1 VWFHVLFYLAVG 12
 œ
 The cDNA library was constructed from polyA RNA of cells
 the amino acid sequence of human Fc receptor I from CDNA clone p90 (see X07372) isolated using
 including humans.
 Seed B;
 374 AA;
 100.0%;
ilarity 100.0%;
Conservative 0
 97US-0861205
 92US-0983647
88US-0160416
 970S-0861205
 90us-0553759
 89US-0379076
 Protein; 374 AA
 /note= "encoded by
 300
 A novel method for cloning cDNAs from mammalian
 0;
 Score 68; DB 19;
Pred. No. 0.00042;
); Mismatches 0;
 GGG"
 Length 374;
 using a
 0
 in cos
 Gaps
 Š
```

Fc receptor I; FcRI; cell surface antigen; lymphocyte; human;

cloning.

0,

```
adhesion to an antibody-coated substrate. The method is useful for the isolation and cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V63442-63) encoding cell surface
 13-JUL-1989;
13-JUL-1990;
21-MAY-1997;
 from a single patient undergoing extracorporeal interleukin-2 induction therapy. Clones p135 (see V63456) and p90 (see X07372) were also obtained. A novel method for cloning cDNAs from mammalian expression libraries is based on transient expression of an antigen in eukaryotic cells and selection of cells expressing the antigen by adhesion to an antibody-coated substrate. The method is useful for
 This is the amino acid sequence of human Fc receptor I (FcRI), as deduced from cDNA clone p98/X2 (see X07373) isolated using a rapid immunoselection cloning method from a cDNA library expressed in COS
 Example 10; Column 53-54; 75pp; English.
 New cloning vector and polylinker - based on existing sequences freefficient cloning and expression of mammalian cDNA(s), especially human lymphocyte antigenic sequences
 WPI; 1998-609251/51.
N-PSDB; X07373.
 Aruffo A,
 03-NOV-1998
 US5830731-A
 Misc-difference
 (GEHO) GEN HOSPITAL
 01-DEC-1992;
 21-MAY-1997;
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 25-FEB-1988;
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Homo sapiens
 Misc-difference
 The cDNA library was constructed from polyA RNA of cells
 88US-0160416.
89US-0379076.
90US-0553759.
97US-0861205.
 /note=
 /note=
 /note=
1.77
 92US-0983647
 97US-0861205
 /note=
256
 /note=
176
 /note=
171
 /note=
159
 /note=
141
 /note=
103
 Location/Qualifiers
 /note=
 note=
 note=
 'note= "encoded by
 note=
 CORP
 "encoded by
 "encoded
 "encoded
 "encoded
 "encoded
 "encoded by
 "encoded
 "encoded
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 "encoded
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 bу
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 Λά
 bу
 bу
 bу
 GTC"
 GGG'
 ATG"
 AAC'
 CCC"
 CTG"
 GGC"
 CAA"
 TCC T
 AAT"
 GAC"
 CTC"
```

for

from mammalian

lymphocytes

(see W80440-55).

The

l surface isolated

/note=

"encoded by

```
RESULT
Y96134
ID YS
XX
AC YS
XX
 В
 γQ
 Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 12; Conservative (
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 panning; immunodiagnosis; diagnosis; immunotherapy; gene th immune disorder; infection; asthma; immune-complex disease; amyloidosis; multiple scierosis; parasitic disease; Fc rece
 Misc-difference
 Key
 Homo
 Macrophage; FcRI; cell surface antigen; human; immunoselection;
 Human macrophage-specific FcRI
 19-DEC-2000
 Y96134;
 Y96134 standard;
 the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and immunotherapeutic applications, including the diagnosts and treatment of immune-mediated infections, diseases, and disorders.
 Sequence
 genes can be expressed in a prokaryotic or eukaryotic host cells to produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V63444) which allow the generalization of very large mammalian expression libraries who
 animals,
 289 vwfhvlfylavg 300
 1 VWFHVLFYLAVG 12
 -difference
 sapiens.
 including
 374 AA;
 (first entry)
 /note=
 /note=
9 51
 /note=
117
 /note=
116
 /note=
 /note=
123
 /note=
122
 /note=
56
 82
 64
 Location/Qualifiers
 note=
 /note=
 note=
 /note=
 /note=
 /note=
 'note=
 Protein; 374
 humans
 encoded.
 "encoded
 "encoded
 "encoded
 "encoded
 "encoded
 "encoded
 "encoded
 "encoded
 "encoded
 "encoded
 "encoded
 "encoded
 "encoded by
 "encoded by
 "encoded
 Score 68;
Pred. No.
 A
 bу
 by.
 уd
 bу
 bу
 bу
 bу
 bу
 bу
 γď
 bу
 уď
 γď
 уd
 Mismatches
 CAT
 ACC"
 ccc,
 GCA"
 GAA"
 CAA"
 CAG"
 CAC"
 ACC"
 TTC"
 .000
 TCC"
 cgg.
 ACC"
 GCG"
 DB 19;
0.00042;
 0
 Length 374;
 Indels
 and disorders of
 gene therapy;
 0;
 Gaps
```

Misc-difference

/note=

"encoded

bу

```
The present sequence is that of a human macrophage specific FoRI, as deduced from cDNA clone pl35 (see A50592), which was isolated from a cDNA library expressed in COS cells using a novel method of the invention designed to isolate CSA nucleic acids. The method is
 Isolated nucleic acid molecule encoding the CD19 cell surface antigen, useful for immunodiagnosis and immunotherapy of immune-mediated infections or disorders, e.g. asthma, immune-complex disease, parasitic
 surfaces other than macrophages, is useful protein of FCRI and a receptor ligand will the potencies of antibodies in therapy.
 immune-complex disease, amyloidosis, parasitic diseases or multiple sclerosis. FCRI is a high affinity receptor for the Fc portion of IGG, normally located on the cell surfaces of macrophages. The ability to interfere with such bonding, or to cause it to occur on surfaces other than macrophages, is useful in therapy. A fusion protein of FCRI and a receptor ligand will be helpful to increase
 IgG, normal
ability to
 animals, including humans.
 treatment of immune-mediated infections,
 proteins they encode, are useful for immunodiagnostic and immunotherapeutic applications, including the diagnosis and
 physical selection of cells expressing the antigen by adhesion to (panning on) an antibody-coated substrate such as a culture dish. CSA nucleic acids isolated by the method of the invention, and the
 based upon transient expression of a CSA in eukaryotic cells and
 Example 10; Column 53-55; 75pp; English
 (GEHO) GEN HOSPITAL CORP
 13-JUL-1989;
23-MAR-1990;
 01'-DEC-1992;
 28-OCT-1998;
 29-AUG-2000
 US6111093-A.
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 Misc-difference
 ?5-FEB-1988;
 2000-586382/55.
DB; A50592.
 92US-0983647.
88US-0160416.
89US-0379076.
90US-0498809.
 98US-0181612.
 9008-0553759
 /note=
216
 /note=
213
 /note=
332
 /note=
 /note-
305
 /note=
268
 /note=
220
 /note-
 8
 "encoded by GGT
 "encoded
 "encoded
 "encoded
 "encoded by
 "encoded by
 "encoded
 "encoded
 These disorders include asthma,
 уd
 ьy
 уd
 γď
 bу
 GTG"
 AAT"
 CGT
 TTC"
 CAA"
 AAT"
 diseases, and disorders in
 The method is
 ÷.
```

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RESULT 10
Y96226
 밁
 Query Match
Best Local
 Human; high affinity Fc receptor; FcgammaRI; immunoglobu infection; immune response; CD64; monocyte; macrophage; eosinophil; HIV; IGG; immunosuppressive; antirhemmatic; antiinflammatory; AIDS; SLE; multiple myeloma; rheumato
 compositions containing recombinant proteins may be used to treat or prevent autoimmune diseases, allergies or tumours, especially AIDS, rheumatoid arthritis or MM.
 receptor consist only of the extracellular portion of the receptor and are not glycosylated i.e. they do not have transmembrane domains or signal peptides. The recombinant proteins may be used in immunoassays to determine the immune status of patients with chronic diseases of the
 monocytes and macrophages, but expression can also be induced on neutrophils and eosinophils. Upon Fc receptor activation, immune effector pathways are activated, leading to immune response. The present sequence may be modified to produce recombinant versions. The recombinant Fc
 FcgammaRI is also known as CD64. Fc receptors play an important role in defending the body against infections. First, pathogens are opsonised by serum immunoglobulins. The resulting complex then binds
 Sequence
 immune system, e.g. AIDS, systemic lupus erythematosus (SLE), multiple myeloma (MM), or rheumatoid arthritis. In addition, pharmaceutical
 cells expressing Fc receptors. FcgammaRI molecules are expressed by
 The present sequence is the human high affinity Fc receptor, FcgammaRI.
 Disclosure;
 Novel Fc receptor lacking transmembrane domains, a signal peptide, glycosylation, useful for diagnosing and treating immune disorders
 03-DEC-1998;
 07-JUN-2000
 EP1006183-A1
 Homo sapiens
 systemic lupus
 11-SEP-2000
 Y96226;
 Y96226 standard;
 (PLAC) MAX PLANCK GES FOERDERUNG WISSENSCHAFTEN
 289
 2000-367968/32
 374 AA;
 Page 26-28;
 (first entry)
 98EP-0122969
 98EP-0122969
 erythematosus;
 Protein; 374
 300
 Fc
 100.0%;
 60pp; English.
 Score 68;
Pred. No.
 tumour.
 FcgammaRI.
0.00042;
thes 0;
 В
 immunoglobulin;
 rheumatoid arthritis;
 Length 374;
 neutrophil; cytostatic;
 ៩
```

Вb

1 VWFHVLFYLAVG

300 12 l Similarity

Conservative

0

Indels

0;

0

Query Match
Best Local Similarity
Matches 12; Conser

Conservative

0;

Mismatches

100.0%;

Score 68; I Pred. No. 0.

DB 21; ).00042; es 0;

Length 374; Indels

0,

1 VWFHVLFYLAVG

Sequence

B43683 ID B4 XX

B43683 standard; Protein;

399

A

B43683;

밁

314 vwfhvlfylavg 325

```
Query Match
Best Local (
 Matches
 vasotropic; antipsoriatic and antianglogenic. The polynucleotides and polypeptides can be used for preventing, treating or ameliorating medical conditions and diagnosing pathological conditions. Polynucleotides, polypeptides, antibodies, agonists and antagonists from the present invention may be used to treat immune disorders by activating or inhibiting the proliferation, differentiation or mobilisation of immune cells, to treat disorders of haematopoietic cells, autoimmune disorders, allergic reactions, graft versus host disease and organ rejection, cancers, cardiovascular disorders, neurological disease and bacterial or manual treations.
 Sequence
 antagonists may be also be B44240 represent sequences
 C77607 to C78448 encode the human cancer associated proteins given in B43398 to B44239. The proteins can have activities based on the tissu and cells the genes are expressed in. Example of activities include:
 Claim 11; Page 1739-1740; 2352pp; English.
 viral infections.
 antithyroid; antiallergic; antibacterial; antiviral; dermatological;
neuroprotective; cardiant; thrombolytic; coagulant; nootropic;
 cytostatic; proliferative;
 Novel isolated nucleic acids comprising sequences encoding peptides useful for treating or diagnosing e.g. cancer -
 invention
 antiasthmatic; antirheumatic; antiarthritic; antiinflammatory
 WPI; 2000-587533/55
 Rosen CA,
 (HUMA-)
 12-MAR-1999;
 08-MAR-2000; 2000WO-US05882
 W0200055350-A1
 Homo sapiens.
 neurological
 vasotropic; antipsoriatic; antiangiogenic; gene therapy; inflammation; immune disorder; haematopoietic cell disorder; autoimmune disorder; allergic reaction; graft versus host disease; organ rejection;
 diagnosis; cytostatic;
antidiabetic; antiasth
 Human; cancer associated gene; cancer antigen; detection; cancer;
 Human cancer
 08-FEB-2001
 haemostatic;
 dermatological;
 antiinflammatory;
 Local Similarity 100 hes 12; Conservative
VWFHVLFYLAVG 12
 HUMAN GENOME
 399 AA;
 Ruben SM
 thrombolytic; cardiovascular disorder; infection; disease; drug screening.
 disease;
 associated protein sequence SEQ ID NO:1128
 (first entry)
 antiasthmatic; antirheumatic; antiarthritic; antiviral;
vry; antithyroid; antiallergic; antibacterial; cardiant;
 99US-0124270.
 s. The peptides, nucleotides, antibodies, agonists and be also be used in drug screens. C78449 to C78457 and
 neuroprotective; thrombolytic; coagulant; nootropic;
 100.0%;
 proteins can have activities based on the tissues
 proliferative;
 used
 vulnerary;
 0;
 Score 68; DB 21;
Pred. No. 0.00045;
); Mismatches 0;
 in drug screens. C78449 in the exemplification
 immunomodulator; antidiabetic;
 vulnerary; immunomodulator;
 Length 399;
 Indels
 0,
 Gaps
 0;
```

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XAXAX
 RESULT 13
W86195
 Д
 Š
 RESULT 12
R12428
 Query Match
Best Local Similarity
Matches 10; Conserv
 ΧŸ
 10-MAY-1999
 W86195
 W86195 standard; Protein; 377
 Sequence
 regulating antibody production in vivo and in plasmaphoresis to remove immune complexes or pathological antibodies. Rabbit IgG bound to cells transfected with the chimeric Fc(gamma) receptor
 This chimeric sequence is one example of a hybrid Fc receptor of the invention. The hybrid Fc receptor retains the Ig binding functions of the constituent receptor types. It can be used for
 See also Q12078.
 Claim 10;
 antibody prodn.
 New immunoglobulin binding hybrid Fc receptor molecules - used to control auto-immune diseases and allergic reactions and to regulate
 25-OCT-1989;
 N-PSDB; Q12079.
 Hogarth PM,
 (UYME-) UNIV MELBOURNE
 25-OCT-1990;
 16-MAY-1991
 W09106570-A
 Region
 R12428;
 R12428 standard; Protein; 410 AA.
 304 vwfhilfylsvg
 Key
 Immunoglobulin-binding molecule; Fc(gamma)RI; Fc(gamma)RII.
 Hybrid Fc(gamma)RII/I receptor.
 14-AUG-1991
 Region
 1 VWFHVLFYLAVG 12
 1991-164135/22
 fig 3B; 80pp; English
 410 AA;
 Conservative
 (first entry)
 Hulett
 (first entry)
 89AU-0007045
 90WO-AU00513
 /label= Fc(gamma)RI D3,
 Location/Qualifiers
 315
 'label= Fc(gamma)RII domains
 Ð,
 94.1%;
 cytoplasmic domains
 Ierino FL,
 Score 64;
Pred. No.
 AA
 Mismatches
 McKenzie IFC,
 0.0022;
 DB 12;
 transmembrane and
 d1 and
 Length 410;
 Indels
 Osman
 z
 0,
 Gaps .
```

Human Fc receptor I

```
RESULT 14
G50156
 Query Match
Best Local Similarity
Watches 9; Conserva
 This polypeptide human Fc receptor I (FCRI). FCRI cDNAs (see W81213) were isolated using a rapid immunoselection cloning method from a cDNA library expressed in COS cells. The cDNA library was constructed from polyA RNA of cells from a single patient undergoing extrcorporeal interleukin-2 induction therapy. DNA sequence analysis revealed that the cDNAs encoded type I integral membrane proteins with 3 extracellular immunoglobulin domains. The invention provides a novel method for cloning cDNAs from mammalian expression libraries. This is based on transient expression of an antigen in eukaryotic cells and physical selection of cells expressing the antigen by adhesion to an antibody-coated substrate. The method is useful for the isolation and antibody-coated substrate. The method is useful for the isolation
 Allen J, A
Oquendo C,
 25-FEB-1988;
13-JUL-1989;
23-MAR-1990;
13-JUL-1990;
 15-DEC-1998.
 Key
Protein
18-OCT-2000
 G50156;
 G50156 standard;
 cDNA encoding human CD40 antigen - useful for cloning cDNA encoding cell surface antigens, constructing cDNA libraries, expression vectors for expression in eukaryotic cells or their fragments
 01-DEC-1992;
 07-JUN-1995;
 Sequence
 and molecular cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. CD40 cDNA (see V81198) is specifically claimed.
 Example 10; Column 55-56; 79pp; English.
 WPI; 1999-069813/06.
 (GEHO.) GEN HOSPITAL CORP
 US5849898-A.
 Homo sapiens
 Fc receptor I;
 1 VWFHVLFYLA 10
|||:|||||
| 292 vwfnvlfyla 301
 Amiot M,
Seed B,
 377 AA;
 Conservative
(first entry)
 92US-0983647.

8BUS-0160416.

89US-0379076.

89US-0498809.

90US-0453759.
 CD36; cell surface antigen; human; cDNA library.
 95US-0485447
 95US-0485447
 Location/Qualifiers
1..377
 /note=
 Protein;
 Aruffo A,
Simmons
 75.0%;
 "this sequence contains a considerable number of differences from the sequence
 deduced from the DNA sequence given in the specification"
 243 AA.
 Score 51; DB 20; Pred. No. 0.36; 1; Mismatches
 D,
 Camerini D, I
 Lauffer L;
c I, Stengelin
 0;
 Length 377;
 Indels
 ŝ
 0;
 Gaps
 0
 18-JUN-1999;
18-JUN-1999;
18-JUN-1999;
18-JUN-1999;
 Protein identification; signal transduction pathway; metabolic pathway; hybridisation assay; genetic mapping; gene expression control; promoter; termination sequence.
 Arabidopsis thaliana.
 25-FEB-1999
05-MAR-1999
 25-FEB-2000;
 Arabidopsis thaliana protein fragment SEQ ID NO: 63527
 18-JUN-
 .8-JUN-1999
 8-JUN-
 1999
 2000EP-0301439
 99US-0139457.
99US-0139458.
99US-0139459.
99US-0139460.
 99US-
 99US-0139454
99US-0139455
 99US-01
 99US-0139492
 Su66
 99US-013007
 99US-0121825
 13494
 26785
```

SXCCCCCCCCCCCCXXXXTTTTXXXX

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99US-0139817. 99US-0139899. 99US-0140353. 99US-0140354. 99US-0140695.

99US-0140823 99US-0140991

```
RESULT
G50155
 Db
 γ
 Query Match
Best Local S
Matches
 28-OCT-1999;
28-OCT-1999;
28-OCT-1999;
29-OCT-1999;
 Protein identification; signal transduction pathway; metabolic pathway; hybridisation assay; genetic mapping; gene expression control; promoter; termination sequence.
 26-OCT-1999;
26-OCT-1999;
26-OCT-1999;
 Arabidopsis
 G50155;
 G50155
 18-OCT-2000
 1 VWEHVLFYLAV 11 :|||:||| : 219 iwfhilfyffi 229
 25-OCT-1999,
25-OCT-1999;
 22-OCT-1999
 21-OCT-1999
 05-OCT-1
 08-OCT
 14-OCT-1999
 L4-OCT-1999
L4-OCT-1999
 .4 -OCT
 2-OCT-1999;
2-OCT-1999;
 15
 standard; Protein; 257 AA
 Similarity 54.96; Conservative
 thaliana
 (first entry)
 99US-0160741
99US-0160767
99US-0160770
99US-0160814
99US-0160815
99US-0160815
99US-0160880
99US-0161404
99US-0161406
99US-0161406
99US-0161359
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99US-0161351
99US-0161352
99US-0161352
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99US-0161361
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990S-0155486
990S-0155659
990S-0156428
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990S-0157117
990S-0157127
990S-0157733
990S-0158232
990S-0158232
990S-0158233
990S-0159233
990S-0159233
990S-0159233
 99US-0151080.
99US-0151303.
99US-0151438.
99US-0151430.
 protein fragment SEQ ID NO:
 72.1%;
 Score 49; DB 2
Pred. No. 0.49;
3; Mismatches
 21;
 'n
 63526
 Length 243;
 Indels
 .0
 Gaps
```

0;

02-AUG-1999; 03-AUG-1999; 04-AUG-1999; 04-AUG-1999; 05-AUG-1999; 05-AUG-1999;

27-JUL-27-JUL-28-JUL-02-AUG-02-AUG-

990S-014508 990S-014508 990S-014508 990S-014508 990S-014508 990S-014519 990S-014512 990S-014521 990S-014521 990S-014521 990S-014521 990S-014531 990S-014531 990S-014531 990S-014531 990S-014531 990S-014538 990S-014538 990S-014730 990S-014730 990S-014730 990S-014730 990S-014730 990S-014730

19-JUL-1999; 19-JUL-1999; 19-JUL-1999;

990S-0144085.
990S-0144325.
990S-0144331.
990S-0144331.
990S-0144333.
990S-0144333.
990S-0144334.
990S-0144352.

3-JUL-1999

99US-0142920. 99US-0142977. 99US-0143542.

99US-0143624 99US-0144005

99US-0142154. 99US-0142055. 99US-0142390. 99US-0142803.

20-JUL-1999

21-JUL-1999

20-AUG-1999 20-AUG-1999 20-AUG-1999 23-AUG-1999 23-AUG-1999 25-AUG-1999 26-AUG-1999 27-AUG-1999 27-AUG-1999

-1999; -1999;

99US-0147416
99US-0147493
99US-0148171
99US-0148171
99US-0148341
99US-0148341
99US-0149368
99US-0149722
99US-0149722
99US-0149722
99US-0149722
99US-0149723
99US-0149723
99US-0149723
99US-0149723
99US-0149723
99US-0149723

|   | PR            | PR           | PR            | PR:                                          | D. 5              | g ;           | ָר<br>ק        | 7 7           | י<br>ג פ      | ג כ           | PR            | PR            | PR            | .PR           | PR            | PR            | PR           | PR           | PR                                    | ָר ק<br>ק     | PR            | PR            | PR            | PR            | p ;           | ק<br>ק       | 2 5     | בי<br>בי      | PR    | PR    | PR    | PR           | PR           | PR    | PR     | PR :       | PR:          | ָ<br>ק        | PR    | PR            | PR           | PR           | PR                 | ב<br>ב        | PR            | PR            | PR            | בר<br>ג       | PR    | PR            | PR:           | מ<br>מ        | PR            | PR            | PR         | ט<br>ג       | PR           | X;                 | PXX            | В            | XX            | ××    | SO            |   |
|---|---------------|--------------|---------------|----------------------------------------------|-------------------|---------------|----------------|---------------|---------------|---------------|---------------|---------------|---------------|---------------|---------------|---------------|--------------|--------------|---------------------------------------|---------------|---------------|---------------|---------------|---------------|---------------|--------------|---------|---------------|-------|-------|-------|--------------|--------------|-------|--------|------------|--------------|---------------|-------|---------------|--------------|--------------|--------------------|---------------|---------------|---------------|---------------|---------------|-------|---------------|---------------|---------------|---------------|---------------|------------|--------------|--------------|--------------------|----------------|--------------|---------------|-------|---------------|---|
|   | 29-JUN-1999;  | 28-JUN-1999; | 24-JUN-1999;  | 23-JUN-1999:                                 | 23 - JUN 1999.    | 22-JUN-1999   | 21 - TIN-1000. | 18-7111-1000  | 18-TIN-1000:  | 16 TIN 1999;  | 18-JUN-1999;   18-JUN-1999: | 17-JUN-1999;                          | 16-TIN-1999;  | 14-JUN-1999;  | 10-JUN-1999;  | 10-JUN-1999;  | 199           | 190           | 04-TUN-1999; | 1 9 9 P | 3 5           | 199   | 19    | 199   | 21-MAY-1999; | 20-MAY-1999; | 19    | MAY-19 | MAY-199    | 14-MAY-1999: | 14-MAY-1999;  |       | 19            | 06-MAY-1999; | 06-MAY-1999; | 05-MAY-1999;       | 199<br>9      | . 1           |               | -199          | 23-APR-1999;  | -199  | L             |               | 01-APR-1999;  | 19            | -19           | MAR-19     | 09-MAR-1999; | 25-FEB-1999; | į                  | 25-FFR-2000    | 06-SEP-2000. | EP1033405-A2. |       | Arabidopsis t |   |
|   | 99US-0140991. | <u>ٺ</u>     | 9905-0140695. | 99US-0140354                                 | 99115-0140353     | 99175-0139899 | 9905-0139703.  | 00116-0130763 | 9905-0139463. | 9905-0139462. | 9905-0139461. | 99US-0139460. | 99US-0139459. | 9908-0139458. | 99US-0139457. | 99US-0139456. | 9905-0139455 | 99US-0139454 | 9905-0139492                          | 9905-0139452. | 990S-0139119. | 99US-0138847. | 9908-0138540. | 9905-0138094. | 99115-0137724 | 9905-0137503 | ب<br>ا  | 99US-0136782. | 13    | 13    | -013  | -013         | 13           | -013  | -013   | -0134      | 50           | 9905-0134218. | -013  | 99US-0132863. | -013248      | -013248      | 9905-0132485       | 99US-0132407. | 99US-0132048. | 99US-0131449. | 99US-0130891. | 9905-0130449. |       | 99US-0129845. | 99US-0128714. | 99US-0127462. | 99US-0126785. | 99US-0126264. | 9US-012578 | 905-012354   |              | 6000 ER 000 ER 000 | 2000EP-0301439 |              | •             |       | thaliana.     |   |
|   |               |              |               |                                              |                   |               |                |               |               |               |               |               |               |               |               |               | •            |              |                                       |               |               |               |               |               |               |              |         |               |       |       |       |              |              |       |        |            |              |               |       |               |              |              |                    |               |               |               |               |               |       |               |               |               |               |               |            |              |              |                    |                |              |               |       |               |   |
|   |               | •            |               |                                              |                   |               |                |               |               |               |               |               |               |               |               |               |              |              | •                                     |               |               |               |               |               |               |              |         |               |       |       |       |              |              |       |        |            |              |               |       |               |              |              |                    |               |               |               |               |               |       |               |               |               |               |               |            |              |              |                    |                |              |               |       |               |   |
| • |               |              |               |                                              |                   |               |                |               |               |               | •             |               |               |               |               |               |              |              |                                       |               |               |               |               |               |               |              |         |               |       |       |       |              |              |       |        |            |              |               |       |               |              | ,            |                    |               |               |               |               |               |       |               |               | •             |               |               |            |              |              |                    |                |              | •             |       |               |   |
|   |               |              |               |                                              |                   |               |                |               |               |               |               |               |               |               |               |               |              |              |                                       |               |               |               |               |               |               |              |         |               |       |       |       |              |              |       |        |            |              |               |       |               |              |              |                    |               |               |               |               |               |       |               |               |               |               |               |            |              |              |                    |                |              |               |       |               |   |
|   |               |              |               |                                              |                   |               |                |               |               |               |               |               |               |               | ,             |               |              |              |                                       |               |               |               |               |               |               |              |         |               |       |       |       |              |              | ٠.    |        |            |              |               |       |               |              |              |                    |               |               |               | ٠.            |               |       |               |               |               |               |               |            |              |              |                    |                |              |               |       |               |   |
|   |               |              |               |                                              |                   |               |                |               |               |               | •             |               |               |               |               |               |              |              |                                       |               |               |               |               |               | ,             |              |         |               |       |       |       |              |              |       |        |            |              |               |       |               |              |              |                    |               |               |               |               |               |       |               |               |               |               |               |            |              |              |                    |                |              |               |       |               | • |
|   |               |              | v             |                                              |                   |               |                |               |               |               |               |               |               |               |               |               |              |              |                                       |               |               |               |               |               |               |              |         |               |       |       | ·     |              |              |       |        |            |              |               |       |               |              |              |                    |               |               |               |               |               |       |               |               |               |               |               |            |              |              |                    |                |              |               |       |               |   |
|   |               |              |               |                                              |                   |               |                |               |               |               |               |               |               |               |               |               |              |              |                                       |               |               |               |               | ٠             |               |              | •       |               |       |       |       |              |              |       |        |            |              |               |       |               |              |              |                    |               |               |               |               |               |       |               |               |               |               |               |            |              |              |                    |                |              |               |       |               |   |
|   | ,             |              |               |                                              |                   |               |                |               |               |               |               |               |               |               |               |               |              |              |                                       | . •           |               |               | ·4.           | ,             |               |              |         |               | •     |       |       |              |              |       |        |            |              |               |       |               |              |              |                    |               |               |               |               |               |       |               |               |               |               |               |            |              |              |                    |                |              |               |       |               |   |
|   |               |              |               |                                              |                   |               |                |               |               |               |               |               |               |               |               |               |              |              |                                       |               |               |               |               |               |               |              |         |               |       |       |       | ٠            |              |       |        |            |              |               |       |               |              | ,            |                    |               |               |               |               |               |       |               |               |               |               | ,             |            |              |              |                    |                |              |               |       |               |   |
|   |               |              | * **          |                                              |                   |               |                |               |               |               |               |               |               |               |               |               |              |              |                                       |               |               |               |               |               |               |              |         |               |       |       |       |              |              |       |        |            |              |               |       |               |              |              | •                  |               |               |               |               |               |       |               |               |               |               | •             |            |              |              |                    |                |              |               |       |               |   |
|   | PR            | PR.          | PR            | מ ק                                          | , ה<br>ל          | מ פ           | ג              | יל נו<br>א נו | ייי           | 27            | . PR          | PR            | PR            | PR            | PR            | PR            | . PR         | ָ ט          | 200                                   | מי נ          | PR            | . PR          | PR            | PR            | ק ק           | 27.2         | אַל     | 707           | PR    | PR    | PR    | PR           | PR           | PR    | PR     | PR         | ל ק          | אַל           | PR    | PR            | PR           | PR           | י ק<br>ק           | אַק           | PR            | PR            | אָפִי ר       | PR            | PR    | PR            | PR            | PR            | PR            | PR            | PR         | PR           | PR           | מקי                | מים בי         | PR           | PR            | PR    | PR            |   |
|   | 13-           | 10-          | 1             | 014                                          | י<br>ני<br>ני     | ۵ د<br>د د    |                |               | 201           | 25            | 23            | 23-           | 20-           | 20-           | 20-           | 18-           |              |              | ـــــــــــــــــــــــــــــــــــــ | 12.           | 11-           | 10-           | - 60          | 09-           | 0 0           | 06-          | 205     | 04-           | 04-   | 03-   | 02-   | 02-          | 02-          | 28-   | 27-    | 27-        | 27-          | 723-          | 23-   | 23-           | 22-          | 22-          | ) L<br>) L<br>! .! | 21-           | 21-           | 21-           | 20-           | 20-           | 19-   | 1             |               | 19-           | - 1           | 1             | 16-        | 14-          | 13-          | 1                  | 08-JU          | 06-JUI       | 01-           | 01-   | 30-           |   |
|   | P-1           | P-1          | P-1           | <u>'                                    </u> | ָרָרָי<br>מַרָּרָ | יי<br>פנ      | 1              |               | 6.1           | 6-            | G-1           | G-1           | G-1           | G-1           | G-1           | G-1           | G - 1        | ٠,           | ין<br>מינו                            | 1             | G-1           | G-1           | G-1           | G-1           | 4 د           | 9 -          | G-1     | G-1           | AUG-1 | AUG-1 | AUG-1 | AUG-1        | AUG-1        |       |        |            |              |               | ::    | JUL-1         |              |              | 1111               | ::            | Ξ.            | [-1           |               |               | F-3   |               | JUL-1         | 12            | 5             | 돐             | -,         | 15           | 5            | <u>-</u>           | 75             |              | 1 - 1 I       |       | JUN-1         |   |
|   | 999;          | 999;         | 999:          | 999,                                         | 0000              | 0 0           | 9 9 9          | 999           | , 666         | , 666         | 999;          | 999;          | 999;          | 999;          | 999;          | 999:          | 999,         | 999.         | 9999                                  | 999           | 999;          | 999;          | 999:          | 999;          | 999           |              | 999,    | 999;          | 999;  | 999;  | 999;  | 999;         | 999;         | 999;  | 999    | 999:       | 000          | 9999          | 999;  | 999;          | 999;         | 999;         | 0000               | 999;          | 999;          | 999;          | 999;          | 9999;         | 999;  | 999;          | 999;          | 999;          | 999;          | 999;          | 999;       | 999          | 999;         | 999:               | 999;           | 999;         | 999;          | ; 666 | 999;          |   |
|   | 990           | 990          | 990           | 1166                                         | 990               | 000           | 999            | 9 9 9         | 990           | 990           | 990           | 990           | 990           | 990           | 990           | 990           | 066          | 1155         | 997                                   | 990           | . 990         | 99U           | 990           | 990           | 991           | 066          | 990     | 990           | 990   | 99U   | 990   | 990          | 99U          | 99U   | 990    | 066<br>066 | 9911         | 066           | 990   | 99U           | 990          | 990          | 1100               | 990           | 990           | 990           | 066           | 066           | 990   | 990           | 000           | 990           | 990           | 99u           | 990        | 990          | 990          | 990                | 1166<br>1166   | 9908         | 990           | 990   | 99U           |   |
|   | s-015         | S-015        | S-015         | S-015                                        | 21012             | 0.015         | 2 - O L U      | CTO           | S-015         | S-015         | S-014          S-014        | S-014                                 | S-014         | S-014         | S-014         | S-014         | S-014         | 0-014         | S-014        | S-014   | S-014         | S-014 | S-014 | S-014 | S-014        | S-014        | S-014 | S-014  | S-014      | S-014        | S-014         | S-014 | S-014         | S-014        | S-014        | S-014              | S-014         | S-014         | S-014         | S-014         | S-014         | S-014 | S-014         | S-014         | S-014         | S-014         | S-014         | S-014      | S-014        | S-014        | S-014              | S-014          | S-0142390.   | S-014         | S-014 | S-014         |   |
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GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.

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US-08-532-795-23
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US-08-663-310-4
US-08-663-310-11
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| US-09-188-930-144 | US-08-602-359A-33 | US-09-234-613-33  | US-08-933-750C-33 | US-09-370-253-18  | 5200327-17         | US-08-318-193-6   | US-08-318-193-39  | US-08-556-597-168 | US-08-620-717A-9  | US-09-184-223-6   | US-08-590-554A-6  | US-08-485-568A-6  | US-08-989-299-12  | US-08-460-114A-4  | US-08-262-338A-4  | US-08-532-795-27  | US-08-532-795-21  |
| Sequence 144, App | Sequence 33, Appl | Sequence 33, Appl | 33,               | Sequence 18, Appl | Patent No. 5200327 | Sequence 6, Appli | Sequence 39, Appl | Sequence 168, App | Sequence 9, Appli | Sequence 6, Appli | Sequence 6, Appli | Sequence 6, Appli | Sequence 12, Appl | Sequence 4, Appli | Sequence 4, Appli | Sequence 27, Appl | Sequence 21, Appl |

# ALIGNMENTS

| AUTHORS: SOLMARUKI, M. AUTHORS: Goebl, M. | ž | SOURCE: | יש | TOPOLOGY: linear | a | H: 453 | RIS | FO | 201-831-3305 |  | TELECOMMUNICATION INFORMATION: | 051 | - | Y/AGENT I | FILING DATE: 16-AUG-1993 | APPLICATION NUMBER: US 08/107,347 | _ | ⊋ | APPLICATION NUMBER: US/08/439,131A | ATA: | SOFTWARE: PatentIn Release #1.0, Version #1.25 | OPERATING SYSTEM: PC-DOS/MS-DOS | PE: FIO |  | 47 | ĸ | STATE: New Jersey | STREET: ONE CYANAMIA PLAZA | American C | CE ADDRESS: | E SEQUENCES: |  | 3512472 | NVENTION: A DNA | APPLICANT: Bard, Marcin | Lai, | RMATION: | o. 5512472 | Sequence 5, Application US/08439131A |  |
|-------------------------------------------|---|---------|----|------------------|---|--------|-----|----|--------------|--|--------------------------------|-----|---|-----------|--------------------------|-----------------------------------|---|---|------------------------------------|------|------------------------------------------------|---------------------------------|---------|--|----|---|-------------------|----------------------------|------------|-------------|--------------|--|---------|-----------------|-------------------------|------|----------|------------|--------------------------------------|--|
|-------------------------------------------|---|---------|----|------------------|---|--------|-----|----|--------------|--|--------------------------------|-----|---|-----------|--------------------------|-----------------------------------|---|---|------------------------------------|------|------------------------------------------------|---------------------------------|---------|--|----|---|-------------------|----------------------------|------------|-------------|--------------|--|---------|-----------------|-------------------------|------|----------|------------|--------------------------------------|--|

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 NFORMATION FOR SEQ ID NO:
 PUBLICATION INFORMATION
 APPLICATION NUMBER: US 08/107,347
FILING DATE: August 16, 1993
ATTORNEY/AGENT INFORMATION:
NAME: Alan M. Gordon
 SOFTWARE: Word Processor CURRENT APPLICATION DATA:
 TOPOLOGY: 1
 SEQUENCE CHARACTERISTICS:
 TELEPHONE: 201-831-3244
 PRIOR APPLICATION DATA:
 COMPUTER READABLE FORM: MEDIUM TYPE: 3.5" 1.
 NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
ADDRESSEE: American Cyanamid Company
 TITLE OF INVENTION:
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217 WF-ILFFISVG 226
 TELEPHONE: ZU1 3305
 CITY: Wayne STATE: New T
 ITLE OF INVENTION:
 JOURNAL: Molecu
VOLUME: 3
PAGES: 263-273
DATE: 1992
 AUTHORS:
 STRANDEDNESS:
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 DESCRIPTION:
 OPERATING SYSTEM: MS DOS
 COUNTRY:
 TITLE: Fission Yeast stsl+ Gene Encodes a Protein TITLE: Similar to the Chicken Lamin B Receptor JOURNAL: Molecular Biology of the Cell
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 AUTHORS:
 2 WEHVLEYLAVG 12
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 07470
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Patent No. 5525496
DATE: 1992
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
 APPLICATION NUMBER: 08/071,880
FILLING DATE: June 1,193
APPLICATION NUMBER: 07/891,301
FILLING DATE: May 29,1992
ATTORNEY/AGENT INFORMATION:
NAME: Hana Dolezalova
MOLECULE TYPE: peptide ORIGINAL SOURCE;
 TELEPHONE: (415) 324-1677
 CLASSIFICATION: 435
PRIOR APPLICATION DATA:
 COMPUTER READABLE FORM:
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COMPUTER: PC
 SOFTWARE: Wordperfect CURRENT APPLICATION DATA:
 NUMBER OF SEQUENCES: 5
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PC
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 lar Biology of the Cell
 PHILLIPS, MOORE, LEMPIO & FINLEY
5 Sherman Avenue, Suite 6
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 SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
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 TELEPHONE: 703-413-3000
 APPLICANT:
 ATTORNEY/AGENT INFORMATION:
NAME: KELBER, STEVEN B.
REGISTRATION NUMBER: 30,073
 NUMBER OF SEQUENCES:
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 Sequence 4, Application US/08663310
Patent No. 5811273
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 APPLICATION NUMBER: FILING DATE: 29-MA
 COUNTRY: UZIP: 22202
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 STATE:
 STREET:
 69 IWAPVLWWIALG 80
 TOPOLOGY:
 STRANDEDNESS:
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 NAME: KELBER, STEVEN B. REGISTRATION NUMBER: 30,073
 OPERATING SYSTEM: PC-DOS/MS-DOS
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PatentIn Release #1.0, Version #1.30

APPLICATION NUMBER:

23-SEP-1996

US/08/663,:310

CLASSIFICATION:

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APPLICATION NUMBER: WO PCT/JP94/02220
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TELEPHONE: (202)672-5300
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26-DEC-1994
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Qy
 В
 Query Match
Best Local Similarity
5; Conserve
 ; TOPOLOGY: 1; MOLECULE TYPE: US-08-663-310-11
 09-006-491-4
 equence 4, Application US/09006491
 INFORMATION FOR SEQ ID NO:
 ENERAL INFORMATION:
 tent No.
 PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 6-235917
FILING DATE: 05-SEP-1994
 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
 PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
FILING DATE: 23-SEP-1996
 CURRENT APPLICATION DATA:
 SEQUENCE CHARACTERISTICS:
LENGTH: 162 amino acids
 REFERENCE/DOCKET NUMBER: 49
TELECOMMUNICATION INFORMATION:
 PRIOR APPLICATION DATA:
 ATTORNEY/AGENT INFORMATION:
 PRIOR APPLICATION DATA:
 CORRESPONDENCE ADDRESS
 NUMBER OF SEQUENCES:
 APPLICANT:
 PPLICANT:
 TELEFAX: 1-
TELEFAX: 904136
 OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Polocia
 ITLE OF INVENTION:
APPLICATION NUMBER: JP 5
 APPLICATION NUMBER: WO PCT/JP94/02220 FILING DATE: 26-DEC-1994
 FILING DATE
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 CITY: Washington
 70 WWPVLWWIALG 80
 CLASSIFICATION:
 COUNTRY:
 NAME: BENT, Stephen A REGISTRATION NUMBER:
 ADDRESSEE:
 APPLICATION NUMBER:
 FILING DATE:
 APPLICATION NUMBER:
 2 WEHVLFYLAVG 12
 FILING DATE:
 TELEPHONE:
 amino acid
 E: Foley & Lardner
3000 K Street, N.W.,
 USA
 KAJIWARA, Susumu
YOKOYAMA, Akihiro
 KONDO,
 Conservative
 (202)672-5399
 PatentIn Release #1.0, Version #1.30
 Inear
 (202)672-5300
 protein
 IMBER: JP 5-348737
27-DEC-1993
 05-SEP-1994
 Keiji
 No.
 50.0%;
45.5%;
 XANTHOPHYLLS AND THE PROCESS FOR PRODUCING THE
 DNA STRANDS USEFUL FOR THE SYNTHESIS OF
 JP 5-348737
 us 08/663,310
 US/09/006,49:
 JP 6-235917
 5972690ihiko
 29,768
 11:
 Score 34; DE
Pred. No. 47;
 Mismatches
 Suite 500
 Length 162
 0,
 Gaps
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 US-09-006-491-11
 Matches
 Query Match
Best Local Similarity
 equence 11,
 ENERAL INFORMATION:
 NFORMATION FOR SEQ ID NO:
 SEQUENCE CHARACTERISTICS:
LENGTH: 162 amino acids
TYPE: amino acid
 ent No.
 TELEPHONE: (202)672-5300
FORMATION FOR SEQ ID NO:
 PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 00
FILING DATE: 23-SEP-1996
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
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 REFERENCE/DOCKET NUMBER: 49441/109
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TELEPHONE: (202)672-5300
 APPLICATION NUMBER: JP 5-348737 FILING DATE: 27-DEC-1993 GTTORNEY/AGENT INFORMATION:
 FILING DATE: 05-SEP-1
PRIOR APPLICATION DATA:
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 CORRESPONDENCE ADDRESS:
 CURRENT APPLICATION DATA:
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 ITLE OF INVENTION:
 69 IWAPVLWWIALG 80
 TELEFAX:
 NAME: BENT, Stephen A. REGISTRATION NUMBER: 29,768
 APPLICATION NUMBER:
 FILING DATE:
 CLASSIFICATION:
 FILING DATE
 APPLICATION NUMBER:
 COUNTRY:
 STREET: 3000-K ST
CITY: Washington
 TOPOLOGY:
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 APPLICATION NUMBER:
 1 VWFHVLFYLAVG 12
 ADDRESSEE:
 20007-5109
 D.C
 904136
 Application US/09006491
 Conservative
 3000 K Street, N.W., Suite 500
 USA
 (202)672-5399
 (202)672-5399
 YOKOYAMA,
 KONDO, Keiji
 KAJIWARA,
 PatentIn
 linear
 05-SEP-1994
 26-DEC-1994
 50.0%;
 XANTHOPHYLLS AND THE PROCESS FOR PRODUCING THE
 DNA STRANDS USEFUL FOR THE SYNTHESIS OF
 XANTHOPHYLLS
 JP 6-235917
 WO PCT/JP94/02220
 Release #1.0, Version #1.30
 Susumu
 US 08/663,310
 US/09/006,49:
 5972690ihiko
 49441/109
 5; Mismatches
 Score 34; DB
Pred. No. 47;
 DB 2;
 Length 162
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"hhes 5; Conserve
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US-09-335-919-4
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Patent No. 615013
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
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LENGTH: 162 amino acids
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PRIOR APPLICATION DATA:
 SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
 TELECOMMUNICATION INFORMATION: TELEPHONE: (202)672-5300
 ATTORNEY/AGENT INFORMATION:
 PRIOR APPLICATION DATA:
 APPLICANT:
 APPLICANT:
 TITLE OF INVENTION:
 ITLE OF INVENTION:
 70 WWPVLWWIALG 80
 OPERATING SYSTEM:
 TOPOLOGY:
 NAME: BENT, Stephen A. REGISTRATION NUMBER: 29,768
 FILING DATE:
 APPLICATION NUMBER: JP 6-235917
 APPLICATION NUMBER: US/0
FILING DATE: 23-SEP-1996
 TOPOLOGY:
 TELEFAX:
 REFERENCE/DOCKET NUMBER:
 APPLICATION NUMBER:
 CLASSIFICATION:
 FILING DATE
 APPLICATION NUMBER:
 COUNTRY:
 STREET: 3000 K S
CITY: Washington
 2 WEHVLEYLAVG
 ADDRESSEE:
 ENGTH:
 INFORMATION:
 20007-5109
 amino acid
 Application US/09335919
 D.C
 E: Foley & Lardner
 162 amino acids
 KAJIWARA, Susumu
YOKOYAMA, Akihir
 Conservative
 (202)672-5399
 linear
 UMBER: JP 5-348737
27-DEC-1993
 05-SEP-1994
 26-DEC-1994
 12
 Keiji
 PC-DOS/MS-DOS
 50.0%;
 XANTHOPHYLLS
 XANTHOPHYLLS
 DNA STRANDS USEFUL FOR THE SYNTHESIS OF
 US/09/335,919
 Akihiro
 WO PCT/JP94/02220
 US/08/663,310
 6150130ihiko
 49441/109
 5; Mismatches
 Score 34; DB
Pred. No. 47;
 AND THE PROCESS FOR PRODUCING THE
 Length 162;
 Indels
 0;
 Gaps
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B

69 IWAPVLWWIALG 80

1 VWFHVLFYLAVG 12

Query Match
Best:Local Similarity
Matches 5; Conserv

Conservative

50.0%;

Score 34; Pred. No.

Length 162;

Indels

0;

Gaps.

0;

70 WWPVLWWIALG 80

Mismatches

RESULT 11 US-09-335-919-11

SENERAL INFORMATION:

APPLICANT:

KONDO, Keiji

No. 6150130ihiko

tent No.

Application US/09335919

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ρy
 , MOLECULE TYPE: US-09-335-919-11
 Query Match 50.0%;
Best Local Similarity 45.5%;
Matches 5; Conservative
 ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 4944
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
 NFORMATION FOR SEQ ID NO: 11: SEQUENCE CHARACTERISTICS:
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 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
 CORRESPONDENCE ADDRESS:
 PRIOR APPLICATION DATA:
 NUMBER OF SEQUENCES:
 PRIOR APPLICATION DATA:
 TITLE OF INVENTION:
 PRIOR APPLICATION DATA:
 STREET: 3000 K STREET: Washington
 TILE OF INVENTION: DNA STRANDS USEFUL FOR THE SYNTHESIS OF THE OF INVENTION: XANTHOPHYLLS AND THE PROCESS FOR PRODUCING THE
 TLE OF INVENTION:
 TOPOLOGY:
 TYPE: amino acids
 APPLICATION NUMBER: JP 5-348737 FILING DATE: 27-DEC-1993
 FILING DATE:
 COUNTRY:
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 APPLICATION NUMBER: US
 CLASSIFICATION:
 FILING DATE:
 APPLICATION NUMBER: US/09/335,919
 ADDRESSEE:
 APPLICATION NUMBER:
 FILING DATE:
 APPLICATION NUMBER:
2 WFHVLFYLAVG 12
|: ||:::|:|
 20007-5109
 D.C
 3000 K Street, N.W., Suite 500
 · USA
 (202)672-5399
 linear
 protein
 05-SEP-1994
 26-DEC-1994
 XANTHOPHYLLS
 Akihiro
 WO PCT/JP94/02220
 JP 6-235917
 US/08/663,310
 49441/109
 Score 34;
Pred. No.
 Mismatches
 DB
47;
 Length 162;
 Indels
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CORRESPONDENCE

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US-08-118-270-49
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Matches 5; Conserv
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Sequence 49, App.--
Sequence 49, App.--
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 TOPOLOGY: 1
MOLECULE TYPE:
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 REFERENCE/DOCKET NUMBER: 54
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
 PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
FILING DATE: 27-MAY-1992
 CURRENT APPLICATION DATA:
 COMPUTER READABLE FORM:
 CORRESPONDENCE ADDRESS:
 APPLICANT: Schuster, David I.
TITLE OF INVENTION: POLYPEPTIDES OF G-COUPLED PROTEIN
TITLE OF INVENTION: RECEPTORS, AND COMPOSITIONS AND METHODS THEREOF
 NAME: BENT, Stephen A.
 APPLICANT:
 211 VWYHTAFSLVM 221
 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
 CITY: Washington
 STRANDEDNESS
 TELEFAX:
 REGISTRATION NUMBER:
 FILING DATE:
 COUNTRY:
 CLASSIFICATION:
 APPLICATION NUMBER:
 MEDIUM TYPE: Floppy disk
 ADDRESSEE:
 1 VWFHVLFYLAV 11
 133, Application US/08332562A
o. 5985599
 INFORMATION:
 20007-5109
 : _261 amino acids amino acid
 INVENTION:
 D.0
 Application US/08118270
 3000 K Street, N.W., Suite 500
 MCKENZIE, Ian F.C. HOGARTH, Mark P.
 HULETT, Mark D. // FOR IMMUNOGLOBULIN
 (202)672-5399
 50.0%; ilarity 45.5%; Conservative
 BONADONNA, Lisa
 HIBBS, Margaret L
 Murphy,
 PatentIn Release #1.0, Version #1.30
 linear
 protein
 31-OCT-1994
ADDRESS:
 Bernadette M
 Randall B.
 136
 US 07/896,457
 US/08/332,562A
 29,768
 54270/119/GRHA
 Score 34; DB 2; Pred. No. 79;
 Mismatches
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 0;
 Gaps
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ADDRESSEE: BROWDY STREET: 419 Seven CITY: Washington STATE: D.C.

419 Seventh

Suite 300

COUNTRY:

USA

20004

Gaps

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В
 PCT-US93-08528-49
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Conservative
 08-118-270-49
 APPLICATION NUMBER: US/08,
FILING DATE: 09-SEP-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07,
FILING DATE: 10-SEP-1992
ATTORNEY/AGENT INFORMATION:
 quence 49, Application PC/TUS9308528 ENERAL INFORMATION:
 TELEFAX: 202-737-3528
TELEX: 248633
NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
FILING DATE: 09-SEP-1993

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/943,236
FILING DATE: 10-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME:
 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
 TITLE OF INVENTION:
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 MOLECULE TYPE:
 CURRENT APPLICATION DATA:
 APPLICANT:
 TELECOMMUNICATION INFORMATION: TELEPHONE: 202-628-5197
 CURRENT APPLICATION DATA:
 ORRESPONDENCE ADDRESS:
 61 VWYYGLFY 68
 COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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 SOFTWARE:
 COUNTRY: UZIP: 20004
 CITY: Washington STATE: D.C.
 STREET:
 STRANDEDNESS:
 TYPE: amino acid
 SOFTWARE:
 APPLICATION NUMBER:
 ADDRESSEE:
 MBER OF SEQUENCES:
 TOPOLOGY:
 REFERENCE/DOCKET NUMBER:
 NAME: Townsend, Kevin
REGISTRATION NUMBER: (
 1 VWFHVLFY 8
 ENGTH:
 419 Seventh Street, N.W.,
 340 amino acids
 USA
 New York University
VENTION: POLYPEPTIDES OF G-COUPLED PROTEIN
VENTION: RECEPTORS, AND COMPOSITIONS AND METHODS THEREOF
 PatentIn Release #1.0, Version #1.25
 Patentin Release #1.0, Version #1.25
 linear
 BROWDY AND NEIMARK
 Floppy disk
 peptide
 Floppy disk
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 US 07/943,236
 PCT/US93/08528
 US/08/118,270
 34,033
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 Score 34; DB 1;
Pred. No. 1e+02;
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MOLECULE TYPE: peptide
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 Sequence 20, Application US/07937609 Patent No. 5319073
 Patent No.
 NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 4039
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703)836-9300
TELEFAX: (703)683-4109
 GENERAL INFORMATION:
 INFORMATION FOR SEQ ID NO:
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 FILING DATE: 01-APR-1992
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LENGTH: 372 amino acid
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 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
 APPLICANT: WANK, Stephen A.
TITLE OF INVENTION: CHOLECYSTOKININ RECEPTOR-ENCODING DN
 SEQUENCE CHARACTERISTICS
 APPLICATION NUMBER: US 07/928,033 FILING DATE: 11-AUG-1992 ATTORNEY AGENT INFORMATION:
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 TELEPHONE: 202 737-3528
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 CLASSIFICATION:
 61 VWYYGLFY 68
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 COUNTRY: USA
ZIP: 22313-0299
 CITY: Alexandria
 STREET:
 ADDRESSEE:
 STRANDEDNESS:
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 ENGTH:
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 amino acids
 ٧A
 1800 Diagonal Road, Suite 500
 340 amino acids
 Conservative
 linear
 Foley & Lardner
 202-628-5197
 single
 50.0%;
62.5%;
 US 07/831,248
 US/07/937,609
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Дb 

Search completed: June 4, 2001, 12:14:33 Job time: 226 sec

US-07-937-609-20

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Maximum Match 1008
Listing first 45 s
 Maximum
 Minimum
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 OM protein -
 Total number of hits satisfying chosen parameters:
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 Sequence:
 Perfect score:
 Title:
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 Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
 DB
BC
 Score
 seq
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 length:
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 Query
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 Gapop 10.0 ,
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 June
 US-09-284-107-29
 GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.
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pir3:*
pir4:*
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 SUMMARIES
 198801
 Fc-gamma RIIIB-alp
 class V chitin syn
ATPase subunit 6 -
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| 555                | 515                | 483                | 370              | 368                | 366              | 254                | 252                | . 242              | 123               | 82                 | . 776            | 621                | 473               | 407                | 405                |  |
| N                  | N                  | N                  | ٢                | N                  | $\vdash$         | N                  | N                  | N                  | 2                 | N                  | N                | N                  | N                 | N                  | Ν                  |  |
| н83043             | H84013             | T16443             | XNVKUD           | I52863             | XNBYUG           | T20882             | G75485             | E81315             | S63055            | T51909 ·           | 148317           | T48646             | S51256            | н81160             | E81946             |  |
|                    |                    |                    | •                |                    | ٠                |                    |                    |                    |                   |                    |                  |                    |                   |                    |                    |  |
|                    |                    |                    |                  |                    |                  | •                  |                    |                    |                   |                    |                  |                    |                   |                    |                    |  |
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|                    |                    |                    |                  |                    |                  | •                  |                    |                    | .•                |                    |                  |                    |                   |                    | ,                  |  |
|                    |                    |                    |                  |                    |                  | •                  |                    |                    |                   |                    |                  |                    |                   |                    | •                  |  |
|                    |                    |                    |                  |                    |                  | •                  |                    |                    |                   | •                  |                  |                    |                   |                    | •                  |  |
| hypothetical prote | hypothetical prote | hypothetical prote | UDPglucosehexose | myosin heavy chain | UDPglucosehexose | hypothetical prote | hypothetical prote | CDPdiacylglycerol- | probable membrane | related to cytochr | cyclin F - mouse | phytoene dehydroge | probable membrane | hypothetical prote | hypothetical prote |  |

## ALIGNMENTS

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C; Species: Homo sapiens (man)
C; Date: 02-Jul-1996 #sequence\_revision

N; Contains: Fc-gamma (IgG)

C; Accession: I55577; I70303

Fc gamma (IgG) receptor I-B splice form 1 precursor - NAlternate names: CD64

receptor I-B splice form

02-Jul-1996 #text\_change 21-Jan-2000

human

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R;Allen,
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 C; Genetics:
 R;Porges, A.J.; Redecha, P.B.; Doebele, R.; Pan, L.C.; Salmon, J.E.; Kimberly, R.P.
J. Clin: Invest. 90, 2102–2109; 1992
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A; Residues: 1-280 < RES>
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Local Similarity 100.0%; Pred. No. 0.
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 1 VWFHVLFYLAVG 12
J.M.;
 preliminary; translated from GB/EMBL/DDBJ
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 A; Title: Nucleotide sequence of three cDNAs A; Reference number: S03018; MUID:89098339
 R;Allen, J.M.; Seed, B.
Nucleic Acids Res. 16, 11824, 1988
 밁
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 N;Alternate
 F;117-170/Domain:
 A;Cross-references: EMBL:X14355; NID:g31333; PIDN:CAA32536.1; PID:g31334 . A;Note: the authors translated the codon ACT for residue 25 as Ala C:Superfamily: Fc gamma receptor I; immunoglobulin homology C:Keywords: alternative splicing; immunoglobulin receptor; transmembrane protein F;117-170/Domain: immunoglobulin homology <IMM>
 R;Allen, J.M.; Seed,
 A; Title: Isolation and expression of functional A; Reference number: A41357; MUID:89100284
 A; Title: Isolation and
 ;Allen, J.M
cience 243,
 Biol. Chem. 2007 13445 13707, Title: Gene organization of the Reference number: A39878; MUID:9
 A; Title: Nucleotide sequence of three cDNAs Reference number: $03018; MUID:89098339
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 A; Accession:
 ; van de Winkel,
 ; Molecule type: mRNA; Residues: 1-344 <ALL2>
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 Query Match
Best Local
 Species: Homo sapiens (man)
 Residues:
 Molecule type: mRNA
 Accession:
 Accession:
 ccession: S03018
 (289
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 1 VWFHVLFYLAVG
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 Invest.
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 Biol. 13,
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 1-344 <ALL1>
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 170304
 Similarity
 S03019
 Seed, B.
 J.G.J.; Ernst, L.K.; Anderson, 56, 13449-13455, 1991
 90, 2102-2109, 1992
 Conservative
 GB:L03418; NID:g184840;
 2182-2192,
 1989
 300
 12
 GB/EMBL/DDBJ
 100.0%;
 of the human high affinity receptor MUID:91302383
 0;
 Score 68; DB 2;
Pred. No. 0.00019;
 family
 gene products
 NID: g31333; PIDN: CAA32536.1;
 for
 PIDN: AAA36049.1;
 for
 the
 high-affinity
 the
 high-affinity Fc
 human high affinity Fc receptor
 human high affinity Fc receptor
 Length 344
 PIDN:CAA32537.1;
 Chiu,
 PIDN:AAA35678.1; PID:g18 Salmon, J.E.; Kimberly,
 in human mononuclear cells.
 human
 PID:g292169
 Fc receptor complementary
 for
 expression
 21-Jul-2000
 receptor complementary
 IgG,
 Gaps
 PID:g31332
 PID: 931334
 PID:g180279
nberly, R.P.
 FcgammaRI (CD64
 and
 0;
 (Fc
 (FCF
 FC gamma (IGG) receptor high affinity - mouse N;Alternate names: high affinity IgG receptor C;Species: Mus musculus (house mouse) C;Date: 18-Jun-1993 #sequence_revision 18-Nov-1994 #text_change 21-Jan-2000 C;Accession: A46480; A43511
R;OSman, N.; Kozak, C. A.; McKenzie, I.F.; Hogarth, P.M.
J. Immunol. 148, 1570-1575, 1992
 A; Reference number: A46480; A; Accession: A46480
 밁
 A; Status: preliminary
 δÃ
 RESULT\
I48471
 A; Title: Structure and mapping
 A46480
 Дb
 Qy
 RESULT
 Matches
 Best
 Query Match
 303
 Local
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1 VWFHVLFYLAVG 12

10;

Conservative

Similarity

94.1%; 83.3%;

Score Pred.

64; No.

DB 2; 0.0009;

Length

0

Gaps

0

type: DNA

MUID: 92166399

encoding

mouse

high

affinity

FC

gamma

RI

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A;Cross-references: EMBL:X70980; NID:g311748; PIDN:CAA50311.1; C;Superfamily: Fc gamma receptor I; immunoglobulin homology C;Keywords: immunoglobulin receptor F;128-180/Domain: immunoglobulin homology <IMM>
 A;Title: Linkage on chromosome 3 of autoimmune diabetes and defective A;Reference number: I48471; MUID:93242399
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 Fc gamma (IgG) receptor high affinity - mouse
C;Species: Mus musculus (house mouse)
C;Date: 02-Jul-1996 #sequence_revision 02-Jul
 Query Match
Best Local Similarity
Themes 12; Conserv
 R; Prins, J.B.; Todd, J.A.; Rodrigues, N.R.; Ghosh, S.; Science 260, 695-698, 1993
 F;117-170/Domain: immunoglobulin homology <IMM2>F;293-313/Domain: transmembrane #status predicte
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 A; Status: preliminary; translated
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292/Domain: oxfracin: f
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 GB:S57204; NID:g298692;
 300
 12
 100.0%; Score 68; 100.0%; Pred. No.
homology
 0,
 from GB/EMBL/DDBJ
 site; carhob...
 Mismatches
 02-Jul-1996 #text_change
 PIDN: AAD13887.1; PID: 94261587
 0.0002
 DB 1;
 Length 374;
 Hogarth,
 Indels
 (Asn)
 (covalent)
 PID:g311749
 P.M.; Wicker,
 21-Jan-2000
 0,
 Gaps
 Fc receptor
 #status
 transmembran
 0,
 L.S.; G
 predi
```

```
hypothetical protein TM0089 - Thermotoga maritima (C;Species: Thermotoga maritima (C;Species: Thermotoga maritima (C;Date: 11-Jun-1999 #sequence_revision 11-Jun-1999 C;Accession: C72420
 RESULT
C72420
 밁
 멍
 Ş
 F;127-179/Domain:
 A; Molecule type: mRNA
A; Residues: 1-404 <SEA>
 Science 277, 1453-1462
A;Title: The complete
 probable membrane
 A;Cross-references: GB:M31314; NID:g200752; PIDN:AAA40056.1; C;Superfamily: Fc gamma_receptor I; immunoglobulin homology
 J. Immunol. 144, 371-378, 1990 A; Title: Molecular cloning and
 R;Sears,
 A; Note: sequence
 A; Reference number: A43511;
 A; Accession: A43511
 ;Title: The complete genome sequence of Escherichia;Reference number: A64720; MUID:97426617
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 ;Blattner, F.R.;
 ;Status: preliminary
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 Date: 12-Sep-1997 #sequence_revision 17-Sep-1997 #text_change 17-Mar-2000
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 Superfamily: Methanobacterium thermoautotrophicum conserved
 Genetics:
 Species: Escher:
 Matches
 Keywords: immunoglobulin receptor; transmembrane protein
 Accession:
 Matches
 Query Match
Best Local
 284-300/Domain:
 82-98/Domain: transmembrane #status predicted <TM3>
105-121/Domain: transmembrane #status predicted <TM4>
 Molecule type: DNA
 Accession:
 258-274/Domain:
 6-192/Domain: transmembrane
 4-330/Domain: transmembrane
 Rose, D.J.; Mau,
 .66/Domain:
 367
 298
 1 VW-FHVLFYLAVG 12
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 1 VWFHVLFYLAVG
 D.W.; Osman, N.; Tate, ol. 144, 371-378, 1990
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 VWFHILFYLSVG 309
 10;
 transmembrane protein
 C72420
 protein TM0089 - Thermotoga maritima (strain MSB8)
 Similarity 53.17; Conservative
 1453-1462, 1997
 Similarity
Stewart, A.M.;
 transmembrane #status
 extracted from
 Conservative
 transmembrane
 transmembrane
 Plunkett III,
 chia coli
 protein yick - Escherichia
 immunoglobulin homology
 B.;
 12
 379
 64.0%;
53.8%;
 94.1%;
 Shao,
 and expression MUID:90111035
Cotton, M.D.; Pratt, M.S.
 #status predicted <TM1>
 G.; Bloch, C.A.; Perna, Y.
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 #status
 #status predicted <TM5>
 #status predicted
#status predicted
 #status
 Score 43.5; D
Pred. No. 3.8;
4; Mismatches
 2;
 Score 64; DB 2;
Pred. No. 0.0011;
 Mismatches
 predicted
 predicted
 predicted
 0f
 <MMI>
 the mouse high affinity Fc receptor
 8
 <TM2>
 <TM6>
 <8MT>
 2;
 #text_change 21-Jul-2000
 0
 1;
 Length 404
 coli
 Length 394;
 not shown
 N.T.; Burland,
Dodson, R.;
Phillips,
 PIDN: AAC76682.1;
 PID:g200753
 hypothetical protein MTH
R.J.; Haft,
ps, C.A.; Ric
 0,
 Gaps
 Gaps
 V.; Riley,
 NCBIN: 85212,
 PID:g1790091
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 0,
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Hickey
on, D.,
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)
당
 R:Fujiwara, M.; Horiuchi, H.; Ohta, A.; Takagi, M. Blochem. Biophys. Res. Commun. 236, 75-78, 1997 A;Title: A novel fungal gene encoding chitin synth A; Reference number: JC5546; MUID:97366599 A; Accession: JC5546
 Nature 399, 323-329, 1999
A;Title: Evidence for lateral gene transfer between Archaea A;Reference number: A72200; MUID:99287316
A;Accession: C72420
A;Accession: C72420
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A;Residues: 1-1852 <FUJ>
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 C;Comment: This enzyme is a membrane-bound
C;Genetics:
 chitin synthase (EC 2.4.1.16) asmA - Emericella nidulans N;Alternate names: chitin-UDP N-acetylglucosaminyltransferase C;Species: Emericella nidulans, Aspergillus nidulans
 밁
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 submitted to the EMBL Data Library, May 1996 A; Reference number: Z19224 A; Accession: T20102
 ş
 A;Gene: TM0089
C;Superfamily: Thermotoga maritima hypothetical protein TM0089
 C; Genetics:
 A;Cross-references: GB:AE001695; GB:AE000512; NID:g4980569; PIDN:AAD35183.1; A;Experimental source: strain MSB8
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 C;Species: Caenorhabditis elegans
C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 15-Oct-1999
 hypothetical
 RESULT
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 Query Match
Best Local
 Query Match
Best Local
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 15 VWSFVLFYLFAG 26
 2 WEHVLFYLAVG 12
 CESP: C50C10.4
 VWFHVLFYLAVG
 type: DNA
 Similarity 54.96; . Conservative
 Similarity
8; Conser
 protein C50Cl0.4 -
 Conservative
 12
 272/3
 60.3%;
 60.3%;
54.5%;
 Caenorhabditis elegans
 Score 41; DB
Pred. No. 8.9;
 from GB/EMBL/DDBJ
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Pred. No. 4.1;
 Mismatches
 Mismatches
 synthase with a myosin motor-like domain
 2
 2
 #text_change 20-Aug-1999
 4
 Length 340;
 Length 153
 Indels
 and
 Bacteria from
 0
 0,
 Gaps
 Gaps
 catalytic
 O
 0;
 PID: 9498
 genome
```

```
A; Authors: Foulger, D.; Fritz, C.; Fujita, M.; Fujita, Y.; Fuma, S.; Galizzi, P. iech, J.; Harwood, C.R.; Henaut, A.; Hilbert, H.; Holsappel, S.; Hosono, S.; Hu Koetter, P.; Koningstein, G.; Krogh, S.; Kumano, M.; Kurita, K.; Lapidus, A.; I A; Authors: Lauber, J.; Lazarevic, V.; Lee, S.M.; Levine, A.; Liu, H.; Masauda, S.; M.; Ogawa, K.; Ogiwara, A.; Oudega, B.; Park, S.H.; Parro, V.; Pohl, T.M.; F. Rieger, M.; Rivolta, C.; Rocha, E.; Roche, B.; Rose, M.; Sadale, Y.; Sato, T.; A; Authors: Schlackoh, S.; Schroeter, R.; Sooffone, F.; Sekiguchi, J.; Sekowska, akeuchi, M.; Tamakoshi, A.; Tanaka, T.; Terpstra, P.; Tognoni, A.; Tosato, V.; Winters, P.; Wipat, A.; Yamamoto, H.; Yamane, K.; Yasumoto, K.; Yata, K.; Yatuhors: Yoshikawa, H.; Danchin, A.; A; Reference number: A69580; MUID:98044033
A;Cross-references: GB:Z99105; GB:AL009126; NID:g2632457; PIDN:CAB11990.1;
 C; Species: Bacillus C; Date: 05-Dec-1997
 A;Reference number: A69580;
A;Accession: A69747
 В
 C;Species: Chlorella virus PBCV-1
C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999
 C;Keywords: ATP; glycosyltransferase; hexosyltransferase F;102-110/Domain: ATP-binding #status predicted <ATP>
 A;Cross-references: EMBL:U42580;
A;Experimental source: specific)
 submitted to the EMBL Data
 hypothetical protein a70L - Chlor
C; Species: Chlorella virus PBCV-1
 밁
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A;Residues: l-88 <GRA>
 A; Accession: T17560
 Status: preliminary; nucleic acid sequence not shown; translation not shown
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Best Local :
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 Matches
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 Ehrlich, S.D.; Emmerson,
 198 FHVLYYLLAG
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 21 IMFHILFYL 29
 1 VWFHVLFYL 9
 -3 FHVLFYLAVG
 S.; Brouillet,
 ; Ogasawara, N.; Moszer, I:; Albertini, A.M.; Alloni, G.; Azevedo, V.; brouillet, S.; Bruschi, C.V.; Caldwell, B.; Capuano, V.; Carter, N.I., S.D.; Emmerson, P.T.; Entian, K.D.; Errington, J.; Fabret, C.; Ferra
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 207
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Pred. No.
 Score 41;
Pred. No.
 from GB/EMBL/DDBJ
 May 1999
 Mismatches
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 ω
 DB
3.5;
 DB
 2
 2;
 #text_change 15-Oct-1999
 Length. 88;
 Length 1852;
 Indels
 Indels
 0;
 0;
 Gaps
 Carter, N.M.;
, C.; Ferrari,
PID:e1182148
 V.; Uchiyama
 .; Hullo, M.F
 0
 subtilis
 Portetell
 A.; Galle
 Yoshida,
 S.; Mauee
 Lardinois
 A.; Sero
 Scanlon
 Berte
 E.Ch
 C; Date: 05-Dec-1997
 DЪ
 Qy
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 QY
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Query Match
Best Local Similarity
Thes 6; Conserv
 A; Experimental C; Genetics:
 A;Cross-references: GB:AE000463; GB:U00096; NID:ġ2367320;
A;Experimental source: strain K-12, substrain MG1655
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A;Note: the nucleotide sequence was submitted to the EMBL Data Library, October 1993
R;Blattner, F.R.; Plunkett III, G; Bloch, C.A.; Perna, N.T.; Burland, V.; Riley, M.;
A.; Rose, D.J.; Mau, B.; Shao, Y.
 C;Date: 19-May-1994 #sequence_revision 01-Sep-1995 #text_change 24-Sep-1999
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 Science 277, 1453-1462,
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 Nucleic Acids Res.
 R;Plunkett III,
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 Status: preliminary; nucleic acid sequence not shown; translation
 Matches
 Query Match
Best Local S
 le: Analysis of the
 296 WEYLLFFLIV
1 VWFHVLFYLAVG 12
 2 WEHVLFYLAV
 Escherichia coli
 1-421 <BLAT>
 Similarity 6; Conserv
 Escherichia
 G.; Burland,
 Conservative
 S40802;
 305
 11
 3391-3398,
 1997
 Escherichia coli
02; MUID:93347969
 57.4%;
50.0%;
 58.8%;
 coli hypothetical
 V.; Daniels, D.L.; Blattner, 3398, 1993
 Score Pred.
 Score 40;
Pred. No.
 Mismatches
 39;
No.
 genome.
 DB
17;
 DB
24;
 protein (glnA-fdhE intergenic
 2;
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 Length 421
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 DNA sequence of the region
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 F.R.
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 0
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 not shown
 Gaps
 Gaps
 region)
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```

fro

```
hypothetical protein MTH768 - Methanobacterium thermoautotrophicum (strain Delta C; Species: Methanobacterium thermoautotrophicum
 171
 IWFYSLIVIAVG 182
```

R;Smith, C; Accession: F69202 ;Smith, D.R.; Doucette-Stamm, L.A.; Deloughery, C.; Lee, H.; Dubois, J.; Aldredge, T. Qiu, D.; Spadafora, R.; Vicaire, R.; Wang, Y.; Wierzbowski, J.; Gibson, R.; Jiwani, S.; Church, G.M.; Daniels, C.J.; Mao, J.; Rice, P.; Noelling, J.; Reeve, J.N. Bacteriol, 179, G.M.; Daniels, C.J 79, 7135-7155, 1997

#sequence\_revision 05-Dec-1997 #text\_change 22-Oct-1999

H)

A; Accession: F69202 A; Reference number: A; Title: Complete genome sequence A69000; quence of Methanobacterium thermoautotrophicum Delta MUID:98037514

A; Molecule type: DNA A; Residues: 1-84 < MTH> preliminary; nucleic acid sequence not shown; translation

A; Experimental source: C; Genetics: GB:AE000855; GB:AE000666; NID:g2621852; ce: strain Delta H PIDN: AAB85271.1; PID:9262

```
Query Match
Best Local Similarity
Thes 4; Conserve
 submitted to the EMBL Data Library, November 1993 A; Description: Mycobacterium leprae cosmid B1496. A; Reference number: S72693 A; Accession: S72779
 뫄
 MLCL536.34 protein -
 C;Superfamily: formate dehydrogenase focB
 formate dehydrogenase homolog yrhG - Bacillus subtilis
 7: Winters, P.; Wipat, A.; Yamamoto, H.; Yamane, K.; Yasumoto, K.; Yata, K.; Yoshida, N.Authors: Yoshikawa, H.F.; Zumstein, E.; Yoshikawa, H.; Danchin, A.; Title: The complete genome sequence of the Gram-positive bacterium Bacilius subtilis. Reference number: A69580; MUID:98044033
 ubmitted to the EMBL Data Library, September 1997; Reference number: Z16918; Recession, W11777
 Rieger, M.; Rivolta, C.; Rocha, E.; Roche, B.; Rose, M.; Sadaie, Authors: Schleich, S.; Schroeter, R.; Scoffone, F.; Sekiguchi, keuchi, M.; Tamakoshi, A.; Tanaka, T.; Terpstra, P.; Tognoni, A.
 ech, J.; Harwood, C.R.; Henaut, A.; Hilbert, H.; Holsappel, S.; Hosono, S.; Hullo, M.F
oetter, P.; Koningstein, G.; Krogh, S.; Kumano, M.; Kurita, K.; Lapidus, A.; Lardinois
 Species: Mycobacterium leprae Date: 16-Jul-1999 #text_change 17-Nov-2000
 Authors: Lauber,
 Date: 05-Dec-1997 #sequence_revision 05-Dec-1997 #text_change 21-Jul-2000
 Cross-references: EMBL:299125; NID:g2398683; PIDN:CAB16178.1; PID:g2398713 Smith, D.R.; Robison, K.
 Molecule type: DNA
Residues: 1-311 <PAR>
 Status: preliminary; translated from GB/EMBL/DDBJ
 Accession: T11017
 Accession:
 Query Match
Best Local Similarity
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 Matches
 Genetics:
 Cross-references: GB:Z99117; GB:AL009126; NID:g2634966; PIDN:CAB14662.1; PID:g2635166 Experimental source: strain 168
 Molecule type: DNA
Residues: 1-266 <KI
 Status: preliminary; nucleic acid sequence not shown; translation not shown
 ; Ehrlich, 5.1.
190, 249-256,)
 Accession: F6997
 Accession: F69974
 Bron,
 .; Ogawa, K.; Ogiwara,
 189 IWFPIMAFVAIG 200
 1 VWFHVLFYLAVG 12
 4 IWFYIALFLAI 14
 1 VWFHVLFYLAV 11
 F.; Ogasawara, N.; Moszer, I.; Albertini, A.M.; Alloni, G.; Azevedo, V.; Berter
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preliminary
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 Conservative
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 55.9%;
 55.9%;
36.4%;
 A.; Oudega,
 Score 38; DB
Pred. No. 23;
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 5; Mismatches
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 Length 84;
 Indels
 Indels
 0; Gaps
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.; Tosato, V.; Uchiyama
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 0
 0;
 Portetell
 S.; Mauee
 Galle
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Search completed: June 4, 2001, 12:15:47 Job time: 280 sec

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Result
No.
 Title:
Perfect score:
 Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
 Minimum DB seq
Maximum DB seq
 Run on:
 OM protein - protein search, using sw model
 Database
 Scoring table:
 Sequence:
 Total number of hits satisfying chosen parameters:
 Searched:
 Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
 Score
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35.5
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length: 2000000000
 Match Length DB
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 BLOSUM62
Gapop 10.0 , Gapext 0.5
 US-09-284-107-29
68
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 93435 seqs, 34255486 residues
 52
52
52
51
51
51
 1 VWFHVLFYLAVG 12
 919
1972
1978
HH1R_RAT
CIW5_HUMAN
SYM_BUCAI
 SECY_AQUAE
SECY_SYNP7
 MYSG_CHICK
STS1_SCHPO
 GAL7_YEAST
GAL7_KLULA
 FCG1_HUMAN
 SYI_THEMA
MYST_RABIT
 ALFC_PEA
 ALFD_PEA
 FRE7_YEAST
 COX1_ACACA
 UL33_RCMVM
 FDHC_METTE
 CRTI_CERNC
 CIW4_MOUSE
 YH90_SYNY3
 YIHN_ECOLI
 MAUN_METEX
 /NS2_MOUSE
 NL4_YEAST
 G2F_MOUSE
 \D04_HUMAN
 SUMMARIES
 RAT
 _YEAST
 93435
 P12314
 Description
 pisum sativ
pisum sativ
aquifex aeo
 gallus gall
schizosacch
 synechococc
rattus norv
homo sapien
 mus musculu
methylobact
 saccharomyc
rickettsia
 homo sapien
 thermotoga
 acanthamoeb
 rat cytomeg
 kluyveromyc
 saccharomyc
 methanobact
 saccharomyc
 mus musculu
 cercospora
 mus musculu
 rattus norv
 synechocyst
 saccharomyc
 homo sapien
bacillus su
 escherichia
 escherichia
```

| 45          | 44          | 43          | 42          | 41          | <b>4</b> 0  | 39          | 38          | 37                                            | 36          | 3<br>5     | 34          |
|-------------|-------------|-------------|-------------|-------------|-------------|-------------|-------------|-----------------------------------------------|-------------|------------|-------------|
| 34          | 34          | 34          | 34          | 34          | 34          | 34.5        | 34.5        | 34.5                                          | 35          | 35         | 35          |
| 50.0        | 50.0        | 50.0-       | 50.0        | 50.0        | 50.0        | 50.7        | 50.7        | 50.7                                          | 51.5        | 51.5       | 51.5        |
| 280         | 261         | 219         | 162         | 162         | 110         | 374         | 94          | 36                                            | 940         | 580        | 555         |
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| P35839 met  | P08508 mus  |             | Q44262 alc  |             |             | P52407 hev  | w           | -                                             | P57249 buc  | P15362 myc | P48919 can  |
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## ALIGNMENTS

EMBL; X14356; CAA32537.1; -.
EMBL; X14355; CAA32536.1; -.
PIR; S03018; S03018.
PIR; S03019; S03019.
PIR; A41357; A41357.
PIR; B41357; B41357.

HSSP; P12319; 1ALT.
MIM; 146760; -.
InterPro; IPR003006;
Pfam; PF00047; ig; 3.

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 Osman N., Kozak C.A., Parameter and mapping of the gene gamma RI and chromosomal location c. J. Immunol. 148:1570-1575(1992).
 SEQUENCE FROM N.A.
MEDLINE=92166399; PubMed=153167
Osman N., Kozak C.A., McKenzie
 FCG1_MOUSE
P26151;
01-MAY-1992 (
 - + + +
 MEDLINE=90111035; PubMed=2136886;
Sears D.W., Osman N., Tate B., McKenzie I.F.C., Hogarth P.M.;
"Molecular cloning and expression of the mouse high affinity receptor for IgG.";
 _MOUSE
 01-MAY 1992 (Rel. 22, Created)
01-MAY 1992 (Rel. 22, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
HIGH AFFINITY IMMUNOGLOBULIN GAMMA FC RECEPT
RCJ (FCRI) (IGG FC RECEPTOR I).

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 Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rođentia;
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 1 VWFHVLFYLAVG 12
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 h 100.
Similarity 100.
12; Conservative
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144:371-378(1990).
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 PubMed=1531670;

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MEDLINE=99226230; PubMed=10209755
Liu J.Y., Miller P.F., Gosink M.,
"The identification of a new fami
 SEQUENCE FROM N.A.
STRAIN-K12 / MG1655;
MEDLINE-93315143; PubMed-7686882;
MEDLINE-93315143; PubMed-7686882;
Burland V.D., Plunkett G. III, Daniels D.L., Blattner F.R.;
"DNA sequence and analysis of 136 kilobases of the Escherichia genome: organizational symmetry around the origin of replicatio Genomics 16:551-561(1993).
 Escherichia coli.
Bacteria; Proteobacteria;
Escherichia.
 EMBL; M31314; AAA40056.1;
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 l Similarity 83.
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31:1845-1851(1999)
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 Receptor; Transmembrane; Glycoprotein;
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 Plunkett G. III, Burland V.D., Daniels D.L., B! "Analysis of the Escherichia coli genome. III. region from 87.2 to 89.2 minutes."; Nucleic Acids Res. 21:3391-3398(1993).
-!- SUBCELULAR LOCATION: INTEGRAL MEMBRANE PRC-!- SIMILARITY: TO E.COLI YQCE.
-!- SIMILARITY: SOME, TO M.GENITALIUM MG294 ANT
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 Escherichia coli.
Bacteria; Proteobacteria;
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 efflux transporters."
 MEDLINE-99367417; PubMed=10438463;
 SEQUENCE FROM N.A.
 NCBI_TaxID=562;
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 Liu J.Y., Miller P.F.,
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 FUNCTION: INVOLVED IN THE EFFLUX OF SUGARS. THE I MAY BE THE DETOXIFICATION OF NON-METABOLIZABLE SI SUBCELLULAR LOCATION: INTEGRAL MEMBRANE PROTEIN.
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 Chem. 274:22977-22984(1999)
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 This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation -
 Caenorhabditis elegans by comparative Genome Res. 10:703-713(2000).
 SEQUENCE FROM N.A. MEDLINE-20272150;
 Xu S.-H., Gu
Huang G.-Y.,
 MEDLINE-20402571; PubMed-10931946;
 TISSUE-Adrenal
 SEQUENCE FROM N.A.
 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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 'Identification of novel human genes aenorhabditis element
 IR.-M., Han Z.-G., Song H.-D., Peng Y.-D., Huang Q.-H., Ren S.-X., IY.-J., Huang C.-H., Li Y.-B., Jiang C.-L., Fu G., Zhang Q.-H., B.-W., Dai M., Mao Y.-F., Gao G.-F., Rong R., Ye M., Zhou J., S.-H., Gu J., Shi J.-X., Jin W.-R., Zhang C.-K., Wu T.-M., lang G.-Y., Chen Z., Chen M.-D., Chen J.-L., lang G.-Y., Wu T.-M., lang G.-Y., lang G.-Y., Wu T.-M., lang G.-Y., lang G
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 This SWISS-PROT entry is copyright. It is produced through a collaboration
 PROSITE;
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 Duesterhoeft A., Ehrlich S.D., "Sequence of the Bacillus subtilis genome region the lev operon reveals two new extracytoplasmic
 InterPro;
 polymerase sigma factors SigV and Sigz.";
Microbiology 143:2939-2943(1997).
 Sorokin A.,
 Bacillus subtilis.
Bacteria; Firmicutes; Bacillus/Staphylococcus
 YRHG_BACSU
005399;
 BACSU
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 SEQUENCE FROM N.A.
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 EMBL; AF110777; AAF14860.1; EMBL; AF151895; AAD34132.1;
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 Bolotin-Fukuhara M., Buhler J.-M., Francingues-Gaillard M.-C.; Submitted (MAR-1992) to the EMBL/Ge
 ARE1 OR SATZ OK ICROSON CO. Saccharomyces cerevisiae (Baker's yeast).
Saccharomycet, Fungi, Ascomycota; Saccharomycotina; Saccharomyces.
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 "Sterol esterification in yeast: a Science 272:1353-1356(1996).
 MEDLINE=96243137; PubMed=8650549; Yang H., Bard M., Bruner D.A., Gl
 SEQUENCE OF 1-328 FROM N.A.
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STEROL O-ACYLTRANSFERASE 1 (EC 2.3.1.:
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 "Sequence analysis of the genome of the unicellular cyanobacterium Synechocystis sp. strain PCC6803. II. Sequence determination of the entire genome and assignment of potential protein-coding regions.";
 Kaneko T., Sato S., Kotani H., Tanaka A., Asamizu E., Nakamura Y., Miyajima N., Hirosawa M., Sugiura M., Sasamoto S., Kimura T., Hosouchi T., Matsuno A., Muraki A., Nakazaki N., Naruo K., Okumura Shimpo S., Takeuchi C., Wada T., Watanabe A., Yamada M., Yasuda M.
 Synechocystis sp. (strain PCC 6803). Bacteria; Cyanobacteria; Chroococcal NCBI_TaxID=1148;
 DNA
 MEDLINE=97061201; PubMed=8905231; Kaneko T., Sato S., Kotani H., Ta
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P72793;
 Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Rodentia;
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-!- SUBCELLULAR LOCATION: INTEGRAL MEMBRANE PROTEIN (POTENTIAL).
-!- SIMILARITY: BELONGS TO THE UPF0093 FAMILY.
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24063 MW;
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 Last annotation update) PROTEIN SLR1790.
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 Chroococcales; Synechocystis
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 Score 37;
Pred. No.
 PRT;
 137CF30151F75EF2 CRC64;
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 EMBL;
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 or send
 "Rat class III Fc gamma receptor isoforms binding specificity and fail to associate
 "Rat CD16 is defined by a family of class III Fc grequiring co-expression of heteroprotein subunits.
J. Immunol. 146:4352-4361(1991).
 SEQUENCE FROM N.A.
 DOMAIN
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 HSSP;
 EMBL;
 EMBL;
 the European Bioinformatics Institute.
 between.
 This SWISS-PROT entry is copyright. It is produced through a collaboration
 GAMMA. LOW AFFINITY RECEPTOR.
-!- SUBUNIT: MAY FORM MULTISUBUNIT COMPLEX WITH OTHER HETEROPROTEINS
 zeta."
 Sears D.W.;
 Farber D.L., Giorda R., Nettleton
 MEDLINE=93246650; PubMed=8482840;
 STRAIN-SPRAGUE-DAWLEY; TISSUE-Brain;
 "Characterization and expression of an from rat natural killer cells.";
 Zeger D.L.,
 MEDLINE=90239026; PubMed=1692135;
 TISSUE=Natural killer cells;
 Farber D.L.,
 MEDLINE-91250730; PubMed=1710249;
 STRAIN-SPRAGUE-DAWLEY
 DOMAIN
 -!- FUNCTION:
 J. Immunol.
 SEQUENCE FROM N.A.
 SEQUENCE FROM N.A.
 ISOFORM H; HAVE BEEN FOUND. THE SEQUENCE SHOWN IS THAT OF ISOFORM C. THEY MAY BE PRODUCED BY ALTERNATIVE SPLICING, BUT THE EXISTENCE OF A FAMILY OF HIGHLY HOMOLOGOUS AND PRESUMABLY DUPLICATED GENES IS ALSO POSSIBLE. THE EXPRESSION OF THE DIFFERENT ISOFORMS IS DIFFERENTIALLY REGULATED.
TISSUE SPECIFICITY: EXPRESSED ON NATURAL KILLERS CRITES OF THE DIFFERENTIALLY REGULATED.
 EXPRESSION. DOES NOT ASSOCIATE WITH CD3 ZETA.
SUBCELLULAR LOCATION: TYPE I MEMBRANE PROTEIN (POTENTIAL)
ALTERNATIVE PRODUCTS: AT LEAST EIGHT ISOFORMS; ISOFORM A
 , M64368; AAA42049.1;
, M64369; AAA42050.1;
, M64370; AAA42050.1;
, M32062; AAA41148.1;
, L08446; AAA41151.1;
, P12319; IALT.
 SIMILARITY: CONTAINS 2 IMMUNOGLOBULIN-LIKE C2-TYPE DOMAINS
 THIS ASSOCIATION IS REQUIRED FOR EFFICIENT
 an email to license@isb-sib.ch).
 the Swiss Institute of
 37
222
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143
 150:4364-4375(1993)
 Hogarth P.M., Sears D.W.;
 Sears D.W.;
 36
 RECEPTOR FOR THE
 domain;
 Sci. U.S.A. 87:3425-3429(1990).
 (ISOFORM H)
 (ISOFORM A).
 221
241
267
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194
104
187
 (ISOFORMS B; C
 39
 license agreement (See http://www.isb-sib.ch/announce/
 Alternative
 CYTOPLASMIC (POTENTIAL).
IG-LIKE C2-TYPE DOMAIN.
IG-LIKE C2-TYPE DOMAIN.
BY SIMILARITY.
BY SIMILARITY.
 QLARISFLQNGEFVSFHPYNVSYSISN
 SWKNKQLTKVLLFQNGKPVRYYYQSSNFSIPK ->
 POTENTIAL.
CYTOPLASMIC
 EXTRACELLULAR
 REGION RECEPTOR
 LOW AFFINITY IMMUNOGLOBULIN GAMMA FC
 Transmembrane;
 FC REGION OF COMPLEXED IMMUNOGLOBULINS
 Bioinformatics
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 There are no restrictions ng as its content is in
 Trucco M., Kochan J.P.,
 productively with
 differ in IgG subclass-
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 (POTENTIAL).
 Usage
 Glycoprotein; Signal;
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 NAT. NEUTOSCI. 2:422-420(1333)
-i- FUNCTION: VOLTAGE INSENSITIVE, INSTANTANE
-i- FUNCTION: VOLTAGE OUTWARD RECTIFICATION
 -
 "A neuronal two P domain K+ channel stimulated by arachidonic acid polyunsaturated fatty acids."; EMBO J. 17:3297-3308(1998).
 KCNK4 OK INC. (Mouse).
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Eukaryota; Moteria; Rodentia;
 MEDLINE=98292450; PubMed=9628867; Fink M., Lesage F., Duprat F., He
 VARIANT
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 Patel A.J., Honore E., Lesage F., Fink M., Romey G., Lazdunski M., "Inhalational anesthetics activate two-pore-domain background K+
 MEDLINE=99254548; PubMed=10321245;
 01-OCT-2000 (Rel.
01-OCT-2000 (Rel.
POTASSIUM CHANNEL
 CIW4_MOUSE O88454;
 SEQUENCE FROM N.A.
 NCBI_TaxID=10090;
 STIMULATED POTASSIUM
 ACTIVATION
 Lazdunski M.
 217 VWFHAAFCLVM 227
 MISCELLANEOUS: ACTIVATED BY ARACHIDONIC ACID AND OTHER UNSATURAL FATTY ACIDS. NOT AFFECTED BY VOLATILE GENERAL ANAESTHETICS SUCH CHLOROFORM, DIETHYL ETHER, HALOTHANE AND ISOFLURANE. SIMILARITY: BELONGS TO THE TWO PORE DOMAIN FAMILY OF POTASSIUM
 SUBCELLULAR LOCATION: INTEGRAL MEMBRANE PROTEIN (POTENTIAL). ALTERNATIVE PRODUCTS: 2 ISOFORMS; 1 (SHOWN HERE) AND 2/TRAAKT/TRUNCATED; ARE PRODUCED BY ALTERNATIVE SPLICING. TISSUE SPECIFICITY: EXPRESSED IN BRAIN, SPINAL CORD AND EYE.
 POTASSIUM CHANNEL. OUTWARD RECT
EXTERNAL K+ CONCENTRATIONS.
SUBUNIT: HOMODIMER (POTENTIAL)
 1 VWFHVLFYLAV 11
 DETECTED IN HEART,
 Similarity 54.96; Conservative
 (Rel. 40, Last sequence update)
(Rel. 40, Last annotation update)
ANNEL SUBFAMILY K MEMBER 4 (TWIK-RELATED ARACHIDONIC ACID-
OTASSIUM CHANNEL PROTEIN) (TRAAK).
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54.5%;
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 SKELETAL MUSCLE,
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 Score 37; DB Pred. No. 33; 1; Mismatches
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 PRT;
 Heurteaux C., Reyes R.,
 -> R (IN ISOFORM DD4D6A7A3F70ED92
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 01-FEB-1996
01-FEB-1996
01-FEB-1996
 nicotianae phytoene dehydrogenase gene.",
Appl. Environ. Microbiol. 60:2766-2771(1994).
-i- EUNCTION: THIS ENZYME CONVERTS PHYTOENE INTO ZETA-CAROTENE V
INTERMEDIARY OF PHYTOFILUENE BY THE SYMMETRICAL INTRODUCTION
DOUBLE BONDS AT THE C-11 AND C-11' POSITIONS OF PHYTOENE.
 LT 11
_CERNC
 Eukaryota; Fungi; Ascomycota; Pezizomycotina; Dothideomycetes et Chaetothyriomycetes incert Mycosphaerellaceae; mitosporic Mycosphaerella
 VARSPLIC
VARSPLIC
 STRAIN-ATCC
 Ehrenshaft M., Daub M.E.
 MEDLINE=94368091;
 SEQUENCE FROM N.
 Cercospora nicotianae.
 PHYTOENE DEHYDROGENASE
 CRTI_CERNC P48537;
 PDH1.
 SEQUENCE
 CARBOHYD
 DOMAIN
 TRANSMEM
 Glycoprotein;
 Isolation, sequence,
 CARBOHYD
 DOMAIN
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 TRANSMEM
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 PRINTS; PR0133 Ionic channel;
 EMBL; AF056492; MGD; MGI:1298234
 DOMAIN
 TRANSMEM
 236 VWFWILFGLA 245
 Pfam; PF02034; TWIK_channel; 1.
 InterPro; IPR000099;
 or send an email to license@isb-sib.ch).
 [nterPro;
 InterPro; IPR001622
 1 VWFHVLFYLA 10
 SIMILARITY: TO BACTERIAL PHYTOENE DEHYDROGENASES AND TO METHOXYNEUROSPORENE DEHYDROGENASE (CRTD).
 COFACTOR: FAD (PROBABLE).
 PATHWAY: CAROTENOID BIOSYNTHESIS.
 SWISS-PROT entry is copyright. It is produced through a
 Similarity
 PR01333; 2POREKCHANEL. nannel; Transmembrane;
 IPR003280;
 (Rel. 33, Created)
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 18366;
 Conservative
 198
235
256
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 Alternative
 STANDARD;
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 PubMed=8085820;
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 mitosporic Mycosphaerellaceae;
 43051 MW;
 54.4%;
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 Last sequence update)
Last annotation update)
E (EC 1.3.-.) (PHYTOENE
 Score 37;
Pred. No.
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N-LINKED (GLCNAC . . .) (POTENT)
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 CYTOPLASMIC (POTENTIAL):
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 PORE-FORMING (POTENTIAL)
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 Ion transport; Potassium
 478A834B7B7AEC92 CRC64;
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 CG2F_MOUSE S: P51944; Q60797; Q1-QCT-1996 (Rel 01-QCT-1996 (Rel 01-QCT-2000 (Rel
 Submitted (MAR-1995) to the EMBL/GenBank/DDBJ databases.

-- FUNCTION: LIKELY TO BE INVOLVED IN THE CONTROL OF THE CELL CYCOLOGIS SPHASE AND G2 (BY SIMILARITY).

-- SUBCELLULAR LOCATION: NUCLEAR (BY SIMILARITY).

-- SIMILARITY: BELONGS TO THE CYCLIN FAMILY. CYCLIN AB SUBFAMILY.
 NP_BIND
TRANSMEM
 Pfam;
 SEQUENCE FROM N.A.
MEDLINE=95284479; PubMed=7767003;
Obermayr F.O., Sutherland H.F., Kraus B., Frischauf A.-M.;
"Mouse cyclin F maps to a conserved linkage group on mouse chromosome"
 CCNF.
 01-OCT-1996 (Rel. 34, Created)
01-OCT-1996 (Rel. 34, Last sequence up
01-OCT-2000 (Rel. 40, Last annotation
G2/MITOTIC-SPECIFIC CYCLIN F.
 entities
 This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation -
 Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Rodentia;
 MOUSE
 PROSITE; PS00982; PHYTOENE_DH; 1.
Carotenoid biosynthesis; Oxidoreductase; FAD; Flavoprotein;
 InterPro; IPR000553; -.
InterPro; IPR001810; -.
Pfam; PF00646; F-box; 1.
Pfam; PF00134; Cyclin; 1.
PROSITE; PS00292; CYCLINS; 1.
 Croci L.,
Pardi R.,
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 SEQUENCE FROM N.A.
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Cyclin; Cell cycle; Cell
Alternative splicing.
 PROSITE; PS50181; FBOX; 1.
 MGD; MGI:102551;
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 NCBI_TaxID=10090;
 Mus musculus (Mouse)
 Transmembrane
 InterPro;
 EMBL; U03903; AAB86988.1;
 109
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 2 WFHVLFYLAVG 12
 European Bioinformatics Institute. The by non-profit institutions as long
 SIMILARITY:
 247766; CAA87695.1; U20612; AAA62317.1; U20636; AAA63152.1; P30274; IVIN.
 WVSVLIYLLVG 611
 Genome 6:149-150(1995).
 PF02032; Phytoene_dh;
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 EMBL;
 de Antoni A., D Angelo M., Dal Pero F., Sartorello F., Pa
Pallavicini A., Lanfranchi G., Valle G.;
"The DNA sequence of cosmid 14-13b from chromosome XIV of
 01-OCT-1996 (Rel. 34, Creat
01-OCT-1996 (Rel. 34, Last
01-NOV-1997 (Rel. 35, Last
VERY HYPOTHETICAL 15.0 KDA
 SGD;
 Saccharomyces cerevisiae (Baker's yeast).
Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Saccharomyces.
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 Saccharomyces cerevisiae reveals an unusually high number of overlapping open reading frames.";
 NCBI_TaxID=4932;
 P53926;
 CONFLICT
 Hypothetical
TRANSMEM
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 entities requires a
 the European Bioinformatics Institute.
 Yeast 13:261-266(1997).
 MEDLINE=97245296; PubMed=9090055;
 SEQUENCE FROM N.A.
 YNL114C OR N1934
 YNL4_YEAST
 Local
 39 VLFHILKWLSVG 50
 1 VWFHVLFYLAVG
 S71390; CALLIAC.
S0005058; YNLL14C.
Transmembrane
 Z69382; CAA93393.1; -. Z71390; CAA95993.1; -.
 Similarity 58.7; Conservative
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Matches 4
 L7_YEAST PR
GAL7_YEAST STANDARD; PR
GAL7_YEAST STANDARD;
P08431; P04398;
20-MAR-1987 (Rel. 04, Created)
01-OCT-1994 (Rel. 30, Last seque
30-MAY-2000 (Rel. 39, Last anno
 FDHC_METTF
Q50568;
15-JUL-1998
15-JUL-1998
15-JUL-1998
 METTE
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SEQUENCE
 01-OCT-1994 (Rel. 30, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
GALACTOSE-1-PHOSPHATE URIDYLYLTRANSFERASE (ECGAL7 OR YBR018C OR YBR0226.
 TRANSMEM
 MEDLING J. Reeve J.N.;
Nolling J. Reeve J.N.;
"Growth- and substrate-dependent transcription of
"Growth- and substrate-dependent transcription of
 SEQUENCE FROM N.A.
 POTENTIAL FORMATE
 Pfam; PF01226; Form_Nir_trans;
 EMBL; U52681; AAC44819.1;
 or send an email to license@isb-sib.ch).
 the European Bioinformatics Institute.
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 -1- FUNCTION: POTENTIAL FORMATE TRANSPORTER.
-1- SUBCELLULAR LOCATION: INTEGRAL MEMBRANE PROTEIN (PROBABLE)
-1- SIMILARITY: BELONGS TO THE FNT, FAMILY OF TRANSPORTERS (TC.)
 MEDLINE=97158688; PubMed=9006048;
 NCBI_TaxID=145262;
 Methanothermobacter.
 Archaea; Euryarchaeota;
 Methanobacterium thermoformicicum
 Transport;
Saccharomyces cerevisiae
 TRANSMEM
 TRANSMEM
 PROSITE;
 InterPro;
 FRANSMEM
 .207
 Bacteriol. 179:899-908(1997)

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 PS01006; FORMATE_NITRITE_TP_2;
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 FORMATE_NITRITE_TP_1;
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 29070 MW;
 52.9%;
 Methanobacteriales; Methanobacteriaceae;
 (Baker's yeast).
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BOD047347BF7B1A4 CRC64;
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 "Sequence and functional analysis of a 7.2 kb fragment of Saccharomyces cerevisiae chromosome II including GAL7 and new essential open reading frame."; yeast 11:79-83(1995).
 MEDLINE=89131252; PubMed=2851900;
Tajima M., Nogi Y., Fukasawa T.;
"Primary structure of the Saccharomyces cerevisiae GAL7 gene.";
 Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes; Saccharomycetales; Saccharomycetaceae; Saccharomyces.
 ACT_SITE
ACT_SITE
CONFLICT
 MEDLINE=95282516;
 STRAIN=S288C
 NCBI_TaxID=4932;
 Citron B.A.,
 Schaaff-Gerstenschlaeger
 SEQUENCE FROM N.A.
 Yeast 1:67-77(1985).
 SEQUENCE FROM N.A.
 CONFLICT
 EMBL;
 STRAIN=CARLSBERGENSIS
 SEQUENCE OF 1-184 FROM N.A
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 yeast GAL7 gene.";
Nucleic Acids Res.
 MEDLINE-84169499;
 MEDLINE-84185433; PubMed-6715281;
 CONFLICT
 Transferase;
 Pfam; PF01087; GalP_UDP_transf; 1.
PROSITE; PS00117; GAL_P_UDP_TRANSF_I;
 EMBL;
 SEQUENCE OF 1-20 FROM N.A.,
 PIR; S45873; XNBYUG.
 Nucleotide sequence of the
 Sequence of
 InterPro;
 InterPro;
 Bacteriol. 158:269-278(1984).
 URIDYLYLTRANSFERASE FAMILY 1.
 SIMILARITY: BELONGS
 PATHWAY: GALACTOSE METABOLISM.
 CATALYTIC ACTIVITY: UTP +
 UDP-GALACTOSE +
 S0000222;
 X81324;
M12348;
X00215;
 K01752;
 P09148; 1HXQ.
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 IPR000880;
 IPR001937;
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181
 the Saccharomyces GAL
 Donelson J.E.;
 Nucleotidyltransferase;
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57
84
 AAA34628.1;
 CAA57105.1;
AAA34627.1;
 CAA25039.1;
 PubMed=2851900;
 PubMed=7762304;
 PubMed=6324089;
 11:8555-8568(1983)
 179
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266
 PYROPHOSPHATE
 TO THE GALACTOSE-1-PHOSPHATE
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 transcriptional initiation region of
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 PROBABLE.

H -> Y (IN REF. 3

P -> H (IN REF. 3

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 Schindwolf T., Lehnert W.,
 ALPHA-D-GALACTOSE 1-PHOSPHATE
 PROBABLE.
EBA67A8722D19F3B CRC64;
 A (IN REF. I (IN REF.
 region and its transcription
 Galactose metabolism
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Query Match

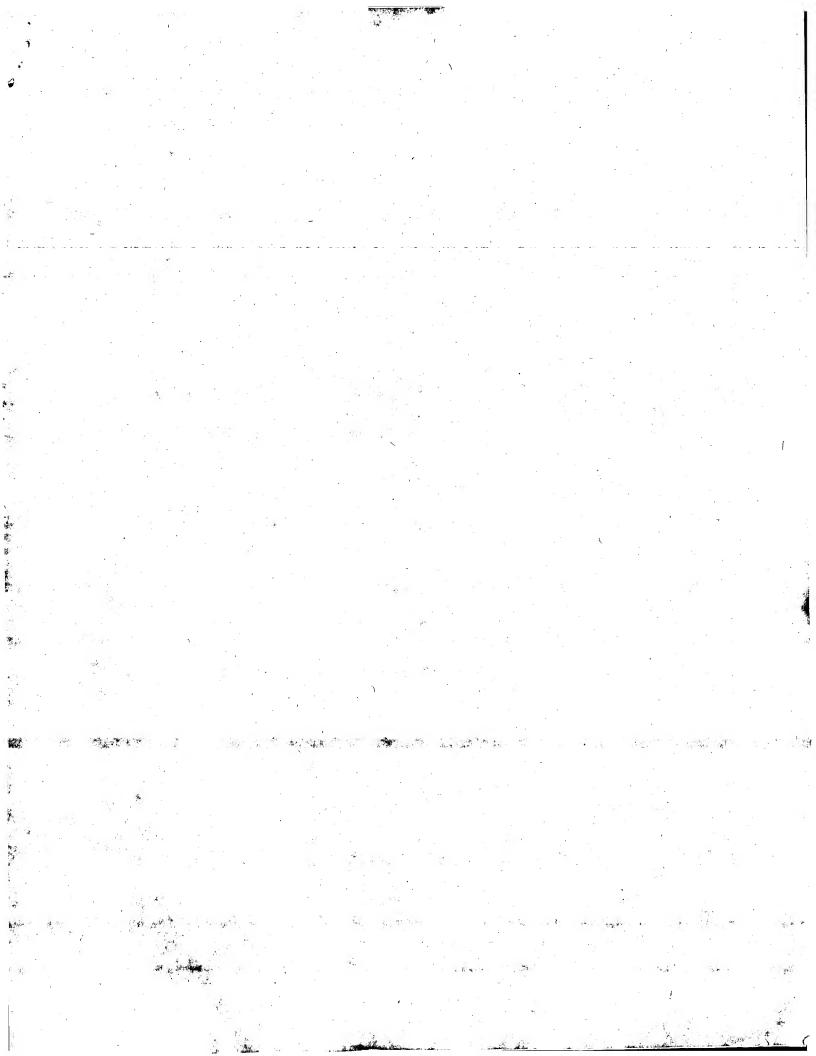
52.98;

Score 36;

DB

٦,

Length 365



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Result
No.
 Database
 Post-processing:
 Minimum DB seq length: 0
Maximum DB seq length: 2000000000
 Run on:
 OM protein
 Total number of hits satisfying chosen parameters:
 Searched:
 Scoring table:
 Sequence:
 Perfect score:
 Title:
 Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
 Score
 38.5
 protein search, using sw model
 Match
 Query
 SPTREMBL_15:*
1: sp_archea:
 Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
 60
 374700 seqs,
 BLOSUM62
Gapop 10.0 ,
 US-09-284-107-29
68
 June
 1 VWFHVLFYLAVG 12
 GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd
 sp_archea:*
 sp_phage:*
 sp_bacteria: *
 sp_unclassified:*
sp_vertebrate:*
sp_virus:*
 sp_organelle:*
 sp_rodent:*
 sp_plant:*
 Length DB
 _invertebrate:*
 4, 2001, 12:23:31; Search time 107.68 Seconds (without alignments)
13.062 Million cell updates/sec
 mhc:*
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 numan:*
 fungi:*
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272
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448
 460
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 Q89405
052498
031428
Q9RQI7
 P89139
Q18731
 Q9P8W4
 Q9WXU4
 Q92638
 Q9M3W5
 SUMMARIES
 374700
 095004
 013281 emericella
Q9p8w4 blumeria gr
Q89405 paramecium
052498 bradyrhizob
 Q9rqi7
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 Description
 Q9m3w5 avena sativ
Q9lnp5 arabidopsis
 P89139 simian
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| 45         | .44        | 43        | 42        | 41        | 40        | 39        | 38        | 37          | 36         | S<br>S     | 34          | 33     | 32     | 31            | 30            | 29          | 28         | 27            | 26         | 25          | 24         | 23          | 22          | 21            | 20          |
|------------|------------|-----------|-----------|-----------|-----------|-----------|-----------|-------------|------------|------------|-------------|--------|--------|---------------|---------------|-------------|------------|---------------|------------|-------------|------------|-------------|-------------|---------------|-------------|
| *          |            |           |           |           |           |           |           |             |            |            |             |        |        |               |               |             |            |               |            |             |            |             |             |               |             |
| 37         | 37         | 37        | 37        | 37        | 37        | 37        | 37        | 38          | 38         | 38         | 38          | 38     | 38     | 38            | 38            | 38          | 38         | 38            | 38         | 38          | 38         | 38          | 38          | 38            | 38          |
| 54.4       | 54.4       | 54.4      | 54.4      | 54.4      | 54.4      | 54.4      | 54.4      | 55.9        | 55.9       | 55.9       | 55.9        | 55.9   | 55.9   | 55.9          | 55.9          | 55.9        | 55.9       | 55.9          | 55.9       | 55.9        | 55.9       | 55.9        | 55.9        | 55.9          | 55.9        |
| 194        | 190        | 148       | 148       | 148       | 148       | 148       | 148       | 1869        | 907        | 497        | 398         | 395    | 393    | 391           | 357           | 354         | 348        | 343           | 314        | 311         | 223        | 197         | 172         | 111           | 84          |
| æ          | Ŋ          | æ         | æ         | œ         | 8         | 8         | 8         | ω           | G          | u          | 10          | 10     | 10     | 10            | 10            | ω           | G          | 10            | <u>ت</u> ر | 2           | N          | N           | σ           | 10            | ۳           |
| 047579     | Q9U5Q4     | Q9T3U4    | Q9T3V8    | Q9T414    | Q9TH29    | Q9TH30    | Q9TH31    | 093809      | OMZA60     | Q9W1D4     | Q9SXX5      | Q9SXX4 | Q9SJU4 | Q9ZU52        | P93565        | Q08417      | Q9W280     | Q9SVJ6        | 058219     | 033142      | 249704     | Q9RKU2      | Q9TTU2      | Q9XEF7        | 026862      |
|            |            |           |           |           |           |           |           |             |            |            |             |        |        |               | •             |             |            |               |            |             |            |             |             |               |             |
|            |            |           |           |           |           |           |           |             |            |            |             |        |        |               |               |             |            |               |            |             |            |             |             |               |             |
|            |            |           |           |           |           |           |           |             |            |            |             |        |        |               |               |             |            |               |            |             |            |             |             |               |             |
| 047579     | Q9u5q4     | Q9t3u4    | Q9t3v8    | Q9t414    | Q9th29    | Q9th30    | Q9th31    | 093809      | Q9vzw0     | Q9w1d4     | Q9sxx5      | Q9sxx4 | Q9sju4 | Q9zu52        | P93565        | . 008417    | Q9w280     | CAS60         | 058219     | 033142      | Q49704     | Q9rku2      | Q9ttu2      | Q9xef7        | 026862      |
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## ALIGNMENTS

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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
 092638
 Kimberly R.P.;
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 1 VWFHVLFYLAVG 12
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cha P.B., Doebele R.,
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 Homo sapiens (Human).
 "Novel Fc gamma
 cells.";
 "Novel Fc gamma receptor I family gene
 J. Clin. Invest. 90:2102-2109(1992)
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 1 VWFHVLFYLAVG
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 EMBL;
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 "Definition of interferon gamma-response elements in a novel human Fc gamma receptor gene (Fc gamma RIb) and characterization of the gene structure ".
 Benech P.D., Sa
Ezekowitz A.B.;
 FC GAMMA RECEPTOR TYPE I (FC CD64.OR FC<GAMMA>RIB.
 Q92495
 Eukaryota; Metazoa;
Mammalia; Eutheria;
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 OMZTO
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 SEQUENCE OF 1-224 FROM N.A
 Submitted (DEC-1992) to the EMBL/GenBank/DDBJ databases
 TISSUE-BLOOD;
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 Eukaryota; Metazoa;
 Homo sapiens (Human)
 Q9MZTO;
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 structure.
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 Bos taurus (Bovine)
 FC GAMMA RECEPTOR I.
 290 VWFHVLFYLAVG 301
 1 VWFHVLFYLAVG 12
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 176:1115-1123(1992)
 Sastry K., Iyer R.R.,
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SEQUENCE
 Zuerner R.L., Harskeer R.A., van de Kemp H., Bal A.E.;

*Sequence analysis of a large ribosomal protein gene operon
Leptospira interrogans."

Submitted (DEC-1998) to the EMBL/GenBank/DDBJ databases.

--- FUNCTION: INVOLVED IN PROTEIN EXPORT.

--- SUBCELLULAR LOCATION: INTEGRAL MEMBRANE PROTEIN.

--- SIMILARITY: BELONGS TO THE SECY/SEC61-ALPHA FAMILY.

EMBL; AF115283; AAD40603.1;
 SEQUENCE FROM N.A.
STRAIN-SEROVAR LAI;
MEDLINE-94041964; PubMed-8409911;
Gravekamp C., van de Kemp H., Franzen M., Carrington D.,
Van Eys G.J., Everard C.O., Hartskeerl R.A., Terpstra W.J
"Detection of seven species of pathogenic leptospires by
sets of primers.";
J. Gen. Microbiol. 139:0-0(0).
 "Molecular cloning and identification of full-length cDNA encoding high affinity Fc.receptor for bovine IgG (Fc gamma RI)."; Vet. Immunol. Immunopathol. 75:151-159(2000).
 01-NOV-1999 (TremBLrel.
01-NOV-1999 (TremBLrel.
01-MAY-2000 (TremBLrel:
PREPROTEIN TRANSLOCASE S
 Q9XD16;
 Q9XD16
 INTERPRO; IPRO02048; INTERPRO; IPRO02208; -PFAM; PF00344; secY; 1
 PRINTS; PR00303; SECYTRNLCASE.
PROSITE; PS00018; EF_HAND; UNKNOWN_1.
PROSITE; PS00095; C5_MTASE_2; UNKNOWN_1.
PROSITE; PS00755; SECY_1; 1.
PROSITE; PS00756; SECY_2; 1.
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 349 AA;
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RA Haft D.H., Hickey E.K., Peterson J.D., Nelson W.C., Ketchum K.A.,
RA McDonald L., Utterback T.R., Malek J.A., Linher K.D., Garrett M.M.,
RA Stewart A.M., Cotton M.D., Pratt M.S., Phillips C.A., Richardson D.,
RA Heidelberg J., Sutton G.G., Fleischmann R.D., Eisen J.A., White O.,
RA Salzberg S.L., Smith H.O., Venter J.C., Fraser C.M.;
RY "Evidence for lateral gene transfer between Archaea and Bacteria from
RY genome sequence of Thermotoga maritima.";
RL Nature 399:323-329(1999).
DR EMBL; AE001695; AAD35183.1; -.
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 01-NOV-1999
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 P89139;
01-MAY-1997
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Zeng L., Godeny E.K.,
 Viruses; ssrNA positive-strand NCBI_TaxID=38143;
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 Gene
 "Molecular characterization of the 3' hemorrhagic fever virus genome.";
J. Virol. 69:2679-2683(1995).
 Godeny E.K., Zeng L.,
 01-MAY-1997
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 Virology
 LARGE GLYCOPROTEIN.
 Mith S.L., Wang
Sequence of the
 15 VWSFVLFYLFAG 26
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8; Conservative
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RESULT Q9WXU4

SEQUENCE FROM N.A. STRAIN-LVR 42-0/M6941;

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CHITIN SYNTHASE D (EC 2.4.1.1
 PFAM; PF00951; Arteri_glycop; PRODOM; PD001151; -; 1 SEQUENCE 278 AA; 31294 MW;
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 Wilson R., Ainscough R., Anderson K., Baynes C., Berks M.,
Bonfield J., Burton J., Connell M., Copsey T., Cooper J., Coulson A
Craxton M., Dear S., Du. Z., Durbin R., Favello A., Fulton L.,
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Jones M., Kershaw J., Kirsten J., Laister N., Latreille P.,
Lightning J., Lloyd C., Mcmurray A., Mortinore B., O'Callaghan M.,
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Smaldon N., Smith A., Sonnhammer B., Staden R., Sulston J.,
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Watson A., Weinstock L., Wilkinson-Sproat J., Wohldman P.,
Watson A., Weinstock L., Wilkinson-Sproat J., Wohldman P.,
 Q18731
 "2.2 Mb of contiguous nucleotide sequence from chromosome III of C. elegans.";
Nature 368:32-38(1994).
EMBL; 27305; CAA96610.1;
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 01-JAN-1998
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 01-JAN-1999
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 IEDLINE=94150718; PubMed=7906398;
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 "The chsD and chsE genes of Aspergillus nidulans and their roles
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 Transferase;
 PFAM; PF00063; myosin_head;
 FRANSMEM
 PRANSMEM
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 novel fungal gene encoding chitin synthase with a myosin motor-like
3 FHYLFYLAVG 12
 GLUCOSAMINYL)] (N+1)
 AB000125; BAA21714.
U62895; AAB05797.1;
 Similarity 70. 7; Conservative
 Biophys.
 family
 IPR002078
 1205
1577
1600
1626
1653
 1843
1852
 Glycosyltransferase; Transmembrane;
 BAA21714.1; -.
 BELONGS
 AA;
 SIGMA54_INTERACT_1; UNKNOWN_1.
 1852
 1225
1597
 1646
1673
 1620
 60.3%;
70.0%;
 Robbins P.W.,
P.J., Rhodes J.
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 Commun.
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ATEAVLSGNL -> GKHRPI (IN STAIN FGSC 4).
 Score 41; DB Pred. No. 99; 1; Mismatches
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 236:75-78(1997).
 CHITIN SYNTHASE FAMILY. SUBFAMILY CLASS
 Bulawa C.E.,
.C., Dodge C.
 Takagi M.;
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 ω,
 Length 1852;
 Iartchouk N.,
 Cell wall;
 (IN STRAIN FGSC 4).
CRC64;
 Indels
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FHVLYYLLAG

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RESULT

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 01-NOV-1996
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052498;
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Q89405;
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MEDLINE-95133167; PubMed-7831789;
Lu Z., Li Y., Zhang Y., Kutish G.F.,
"Analysis of 45 kb of DNA located at
 Zhang Z.G., Gurr S.J.;
Submitted (APR-2000) to the
EMBL; AF189366; AAF04279.2;
SEQUENCE 1867 AA; 207129
 Blumeria graminis.
Eukaryota; Fungi;
NCBI_TaxID=34373;
 Zhang Z.G., Hall A., Person Differential expression
 Hypothetical protein. SEQUENCE 88 AA; 10265 MW;
 3 FHVLFYLAVG
||||:|| |
| 196 FHVLYYLLAG
 STRAIN-CC1;
 CHS2
 CHITIN SYNTHASE.
 Q9P8W4;
 EMBL;
 Virology
 Paramecium bursaria chlorella virus 1
Viruses; dsDNA viruses, no RNA stage;
 HYPOTHETICAL
 STRAIN=CC1;
 SEQUENCE FROM N.A
 SEQUENCE FROM N.A.
 01-OCT-2000
 01-OCT-2000
01-OCT-2000
 NCBI_TaxID=10506;
 21
 Local Similarity
nes 6; Conserv
 13
 1 VWFHVLFYL 9
 12
 Plant Pathol.
 IMFHILFYL
 PBCV-1 genome.";
ogy 206:339-352(1995).
U42580; AAC96438.1; -.
 (TrEMBLrel. 15,) (TrEMBLrel. 15,)
 Conservative
 PRELIMINARY;
 (TrEMBLrel. 01, (TrEMBLrel. 01, (TrEMBLrel. 08,
 PRELIMINARY;
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 Ascomycota;
 1:125-138(2000).
 58.8%;
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 t E., Gurr S.J.;
two chitin synthase
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 Score 40;
Pred. No.
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Pred. No.
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 Rock D.L., Va
 (PBCV-1).
Phycodnaviridae;
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nd of the chlorella
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Matches 6; Conser
 O31428;
O1-JAN-1998
O1-JAN-1998
O1-MAY-2000
 01-JUN-1998
01-JUN-1998
01-JUN-2000
 Azevedo V., Bertero M.G., Bessieres P., Bolotin A., Borchert S.,
Borriss R., Bourster L., Brans A., Brann M., Brighell S.C., Bron S.,
Brouillet S., Bruschi C.V., Caldwell B., Capuano V., Carter N.M.,
Choi S.K., Codani J.J., Connerton I.F., Cummings N.J., Daniel R.A.,
Denizot F., Devine K.M., Dusterhoft A., Ehrlich S.D., Emmerson P.T.,
Entian K.D., Errington J., Fabret C., Ferrari E., Foulger D.,
Fritz C., Fujita M., Fujita Y., Fuma S., Galizzi A., Galleron N.,
Ghim S.Y., Glaser P., Goffeau A., Golightly E.J., Grandi G.,
Guiseppi G., Guy B.J., Haga K., Haiech J., Harwood C.R., Henaut A.,
Hilbert H., Holsappel S., Hosono S., Hullo M.F., Itaya M., Jones L./,
Joris B., Karamata D., Kasahara Y., Klaerr-Blanchard M., Klein C.,
Kobayashi Y., Koetter P., Koningstein G., Krogh S., Kumano M.,
Kurita K., Lapidus A., Lardinois S., Lauber J., Lazarevic V.,
Kurita K., Lapidus A., Lardinois S., Lauber J., Lazarevic V.,
Kurita K., Lapidus A., Lardinois S., Lauber J., Lazarevic V.,
Kurita K., Lapidus A., Lardinois S., Lauber J., Lazarevic V.,
Sekiguchi J., Sekowska A., Seror. ..., Takahashi H., Ta
Sorokin A., Tacconi E., Takagi T., Takahashi H., Ta
Takeuchi M., Tamakoshi A., Tanaka T., Terpstra P., Takeuchi M., Vannier F., Takeuchi M.,
 Kurita K., Lapidus A., Lardinois S., Lauber J., Lazarevic V., Lee S.M., Levine A., Liu H., Masuda S., Mauel C., Medigue C., Medido R.P., Mizuno M., Moestl D., Nakai S., Noback M. Noone D., O'Reilly M., Ogawa K.; Ogiwara A., Oudega B., Park S.H., Park V., Polita M., Portetelle D., Porwoilik S., Prescott A.M., Park S.H., Becker B.U., Kosch K., Parniske M., Muller P.; Submitted (DEC-1997) to the EMBL/GenBank/DDBJ databases. EMBL; AF039306; AAC04824.1; -. INTERPRO; IPR000051; -.
 Bradyrhizobium japonicum.
Bacteria; Proteobacteria;
 Presecan E., Pujic P., Purnelle B., Rapoport G., Rey M., Re
Rieger M., Rivolta C., Rocha E., Roche B., Rose M., Sadaie
Sato T., Scanlan E., Schleich S., Schroeter R., Scoffone F
 Bacillus subtilis.
Bacteria; Firmicutes; Ba
Bacillus/Staphylococcus
NCBI_TaxID=1423;
 031428
 SEQUENCE
 META.
 Kunst F., Ogasawara N., Moszer I., Albertini A.M., Alloni G., Azevedo V., Bertero M.G., Bessieres P., Bolotin A., Borchert
 STRAIN-168
 STRAIN-USDA 110SPC4;
 NCBI_TaxID=375;
 Bradyrhizobium group;
 MEDLINE=98044033;
 SEQUENCE FROM N.A.
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 BDB PROTEIN
 165 WEHVWFY 171
 INTERPRO; IPRO01601;
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 266 AA;
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 PRELIMINARY;
 Sekowska A., Seror S.J., Serror P., Shin B.S.,
Sekowska A., Seror S.J., Serror P., Shin B.S.,
Sekowska A., Seror S.J., Serror P., Shin B.S.,
 PubMed=9384377;
 29164
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 Bacillus/Clostridium group;
us group; Bacillus.
 ria; alpha subdivision;
Bradyrhizobium.
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 Score 40;
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 PRT;
 0318A45D2A1B958C CRC64;
 Mismatches
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 Rhizobiaceae
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 Length 266;
 Rey M., Reynolds S
 Indels
 Vassarotti A.,
 Shin B.S.,
 Noback M.
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 Query Match
Best Local Similarity 60.0
Matches 6; Conservative
 Query Match
Best Local Similarity
Matches 5; Conserv
 STRAIN-168;

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Liu H., Yasumoto K., Takahashi H., Yoshikawa H.;
Haga K., Liu H., Yasumoto K., Takahashi H., Yoshikawa H.;
Haga K., Liu H., Yasumoto K., Takahashi H., Yoshikawa H.;

"Sequence analysis of the 70kb jegion between 17 and 23 degree of
Bacillus subtilis chromosome.";

Submitted (JUL-1997) to the EMBL/GenBank/DDBJ databases.

EMBL; 299105; CAB11990.1;

EMBL; 299105; CAB11990.1;

EMBL; AB006424; BAA33993.1;

INTERPRO; IPR00179;

TOTOCHROME_B_00; UNKNOWN_1.

SEQUENCE 448 AA; 51880 MW; - 3870F3A47E445548 CRC64;
 OPROI7 PRELIMINARY; PRT; 168 AA.
OPROI7;
O1-MAY-2000 (TrEMBLrel. 13, Created)
O1-MAY-2000 (TrEMBLrel. 13, Last sequence update)
O1-JUN-2000 (TrEMBLrel. 14, Last annotation update)
HYPOTHETICAL 18.7 KDA PROTEIN (FRAGMENT)
 STRAIN-EGD;
FSihi H., Cossart P.;
FSihi H., Cossart P.;
"STM applied to Listeria monocytogenes.";
Submitted (OCT-1998) to the EMBL/GenBank/DDBJ databases.
EMBL, AF102170; AAF04741.1;
 STRAIN-168;
Kunst F., Ogasawara N., Yoshikawa H., Danchin A.;
Submitted (NOV-1997) to the EMBL/GenBank/DDBJ databases.
 SEQUENCE FROM N.A.
 Winters'P., Wipat A., Yamamoto H., Yamane K., Yasumoto K., Yata K., Yoshika K., Yoshikawa H.F., Zumstein E., Yoshikawa H., Danchin A.; Toshikawa H.E., Zumstein E., Yoshikawa H., Danchin A.; The complete genome sequence of the gram-positive bacterium Bacillus subtilis.
 Listeria monocytogenes.
Bactleria; Firmicutes; Bacillus/Clostridium group;
Bacillus/Staphylococcus group; Listeria.
NCBI_TaxID=1639;
 SEQUENCE
 Nature 390:249-256(1997).
 INTERPRO; IPRO00566; -. ROSITE; PS00213; LIPOCALIN; UNKNOWN_1.
 296 WEYLLEFLIV 305
 EQUENCE FROM N.A.
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86 IWFPVMAFVAIG 97
 1 VWFHVLFYLAVG 12 :|| |: :: |: |
 2 WFHVLFYLAV-11
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 168 AA; 18718 MW;
 Conservative
 58.8%; Score 40; DB 2; Length 448; 60.0%; Pred. No. 38; Eive 3; Mismatches 1; Indels
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Pred. No. 22;
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 Length 168;
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